



SEQUENCE LISTING

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<120> Nucleotide and Deduced Amino Acid Sequences of the Envelope 1 and Core Genes of Isolates of Hepatitis C Virus and the use of Reagents Derived From These Sequences in Diagnostic Methods and Vaccines

<130> 20264116US2

<140> 09/084,691

<141> 1998-05-26

<150> 08/290,665

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<150> 08/086,428

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<160> 274

<170> PatentIn Ver. 2.1

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<211> 576

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<223> Individual Isolate: DK7

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<213> Homo sapiens

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<213> Homo sapiens

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<212> DNA
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<212> DNA
<213> Homo sapiens

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<212> DNA
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<220>

<223> Individual Isolate: DK1

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<211> 576

<212> DNA

<213> Homo sapiens

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<223> Individual Isolate: HK3

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<212> DNA

<213> Homo sapiens

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<213> Homo sapiens

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<211> 576
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<212> DNA
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<223> Individual Isolate: P10

<400> 18

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<212> DNA

<213> Homo sapiens

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<223> Individual Isolate: S9

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<210> 22
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<212> DNA
<213> Homo sapiens

<220>
<223> Individual Isolate: SW2

<400> 22
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ctggtgttat cgcaagtact ccggatccca caagctgtcg tggacatggt agcgggggcc 480
cactggggag tcctggcggg cttgcatac tattccatgg tggggaaactg ggctaagggtt 540
ttgattgtga tgctactctt tgctggcggtt gacggg 576

<210> 23
<211> 576
<212> DNA
<213> Homo sapiens

<220>
<223> Individual Isolate: T3

<400> 23
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ctagtgggtgt cgcagttgtc ccggatccca caagctgtcg tggacatggt ggcgggggcc 480
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ttgattgtgc tgctactctt tgccggcggtt gatggg 576

<210> 24
<211> 576
<212> DNA
<213> Homo sapiens

<220>
<223> Individual Isolate: T10

<400> 24
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gagggcaact cctcccgctg ctgggttagcg ctcactccca cgctcgccgc caggaacacc 180
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<210> 25

<211> 576
<212> DNA
<213> Homo sapiens

<220>
<223> Individual Isolate: US6

<400> 25
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ctgattgtgt tgctactt tgccggcgtt gacggg 576

<210> 26
<211> 576
<212> DNA
<213> Homo sapiens

<220>
<223> Individual Isolate: T2

<400> 26
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ctggaaata catcccgatg ctggataccg gtcacaccaa acgtggccgt gccgcagccc 180
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attgtcatcc tcttgctggc tgctgggggtg gacgcg 576

<210> 27
<211> 576
<212> DNA
<213> Homo sapiens

<220>
<223> Individual Isolate: T4

<400> 27
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gttgcaccc ttctgctggc cgctggggcg gacgacgc 576

<210> 28
<211> 576
<212> DNA
<213> Homo sapiens

<220>
<223> Individual Isolate: T9

<400> 28
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gttgcaccc ttctgctggc cgctggcggtg gacgacgc 576

<210> 29
<211> 576
<212> DNA
<213> Homo sapiens

<220>
<223> Individual Isolate: US10

<400> 29
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gttgcatcc ttctgctagc cgctggggcg gacgacg 576

<210> 30
<211> 576
<212> DNA
<213> Homo sapiens

<220>
<223> Individual Isolate: DK8

<400> 30
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atcacctggc aactcaccga cgcaatttcac cacccccc gatgcgtccc atgtgagaat 120
gacaatggca ccctgcgtg ctggatacaa gtgacaccta atgtggctgt gaaacaccgc 180
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attgcacatcc tccttcttgt cgcaaggatg gatgc 576

<210> 31
<211> 576
<212> DNA
<213> Homo sapiens

<220>
<223> Individual Isolate: DK11

<400> 31
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gacaatggca ccctgcactg ctggatacaa gtgacaccta atgtggctgt gaaacaccgc 180
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cattggggcg tgggtttgg ctggcctat ttctccatgc agggagcgtg ggcgaaagtc 540
attgcacatcc tccttcttgt agcaggatg gatgc 576

<210> 32
<211> 576
<212> DNA
<213> Homo sapiens

<220>
<223> Individual Isolate: SW3

<400> 32
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cattggggcg tggtgtttgg cttggcctat ttctccatgc aaggagcgtg ggccaaggtc 540
atgcgcattcc tcctgcttgt cgcaaggatg gatgca 576

<210> 33
<211> 576
<212> DNA
<213> Homo sapiens

<220>
<223> Individual Isolate: T8

<400> 33
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cattggggcg tggtgtttgg cttggcctat ttctccatgc aaggagcgtg ggccaaggtc 540
atgcgcattcc tcctccttgt cgcaaggatg gacgca 576

<210> 34
<211> 576
<212> DNA
<213> Homo sapiens

<220>

<223> Individual Isolate: S83

<400> 34

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gtcgtgtcgc cacaacacca tacgttgtc caggaatgca actgttccat ataccggc 360
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atgctcctgg cgtacttggt ggcgcattccg gaagtcatct tggatattgt tacaggaggt 480
cattggggtg taatgtttgg cctcgcttac ttctccatgc agggatcgtg ggcgaaggc 540
atcgttatcc tcctgctgac tgctgggtg gaggcg 576

<210> 35

<211> 576

<212> DNA

<213> Homo sapiens

<220>

<223> Individual Isolate: DK12

<400> 35

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atggtggtag cgacgtcct gcgtctgccc cagacctgtc tcgacataat agctggggcc 480
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gctatcatca tggttatgtt ttcaggagtc gatgc 576

<210> 36

<211> 576

<212> DNA

<213> Homo sapiens

<220>

<223> Individual Isolate: HK10

<400> 36

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gctatcatca tggtatgtt ttcaggggtc gatgcc 576

<210> 37
<211> 576
<212> DNA
<213> Homo sapiens

<220>
<223> Individual Isolate: S2

<400> 37
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attgtgtatg aggccgatga cgttattctg cacacacctg gctgtgtacc ttgtgttcag 120
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<210> 38
<211> 576
<212> DNA
<213> Homo sapiens

<220>
<223> Individual Isolate: S52

<400> 38
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gacggcaata catccatgtg ctggacccca gtgacaccta cggcgtggcagt caggtacgtc 180
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gctattgtca tgattatgtt ttcaggggtc gatgcc 576

<210> 39
<211> 576
<212> DNA
<213> Homo sapiens

<220>
<223> Individual Isolate: S54

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<210> 40
<211> 576
<212> DNA
<213> Homo sapiens

<220>
<223> Individual Isolate: Z4

<400> 40
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<210> 41

<211> 576
<212> DNA
<213> Homo sapiens

<220>
<223> Individual Isolate: Z1

<400> 41
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<210> 42
<211> 576
<212> DNA
<213> Homo sapiens

<220>
<223> Individual Isolate: Z6

<400> 42
gttaactatc gcaatgcctc gggcgtctat cacgtcacca acgactgccccc gaactcgagc 60
atagtgtatg aggccaaaca ccagatctt caccccccag ggtgcttgcc ctgtgtgagg 120
gttggaaatc agtcacgctg ctgggtggcc cttactccca ccgtggcggt gtcttataatc 180
ggtgctccgc ttgactccct ccggagacat gtggacctga tgggggcgc cgctactgt 240
tgctctgccc tctacgttgg agatctgtgc ggtgggtcat tcttgggttgg ccagatgttc 300
tccttccagc cgcgacgcca ctggactacg caggactgca attgttctat ctacgcaggg 360
catatcacgg gccacaggat ggcatggac atgatgtga actggagtcc cacaaccacc 420
ctgcttctcg cccaggtcat gaggatccct agcactctgg tagatctact cgctggaggg 480
cactggggcg tccttggtgg gttggcgtac ttcaagcatgc aagctaattg gccaaagtc 540
atcctggtcc ttttccctttt cgctggagtt gatgcc 576

<210> 43
<211> 576
<212> DNA
<213> Homo sapiens

<220>
<223> Individual Isolate: Z7

<400> 43

gtcaactatc acaatgcctc gggcgtctat cacatcacca acgactgccc gaactcgagc 60
ataatgtatg aggccgaaca ccacatccta cacctccag ggtgcgtacc ctgtgtgagg 120
gaggggaacc agtcacgctg ctgggtggcc cttaactccca ccgtggcggc gccttatatc 180
ggtgcaccgc ttgaatccat ccggagacat gtggacctga tggtaggcgc tgctacagtg 240
tgctccgctc tctacattgg ggacctgtgc ggtggcgtat ttttggttgg tcagatgttt 300
tctttccagc cgcgacgcca ctggactacg caggactgca attgttccat ctatgcgggg 360
cacgttacag gccacagaat ggcatggac atgatgatga actggagtc cacaaccacc 420
ttggtcctcg cccaggttat gaggatccct agcactctgg tggacctact cactggaggg 480
cactgggtta tccttatcgg ggtggcatac ttctgcatac aagctaattg ggccaaggtc 540
attctggtcc ttttcctcta cgctggagtt gatgcc 576

<210> 44

<211> 576

<212> DNA

<213> Homo sapiens

<220>

<223> Individual Isolate: DK13

<400> 44

tacaactatc gcaacagctc gggtgtctac catgtcacca acgattgccc gaactcgagc 60
atagtctatg aaaccgatta ccacatctta caccccccgg gatgcgttcc ttgcgtgagg 120
gaagggaaaca agtctacatg ctgggtgtct ctcaccccca ccgtggctgc gcaacatctg 180
aatgctccgc ttgagtcctt gagacgtcac gtggatctga tggtaggcgg cgccactctc 240
tgctccgccc tctacatcgg agacgtgtt ggggggtgt tcttggtcgg tcaactgttc 300
accttccaac ctcgcccggca ctggaccacc caagactgca attgttccat ctacacagga 360
catatcacag gacacagaat ggcattggac atgatgatga attggagccc cactgcgacg 420
ctggtcctcg cccaaacttat gaggatccca ggccatgg tcgacactgct tgcaggcggc 480
cactggggca ttctggttgg catagcgtac ttctgcatac aagctaattg ggccaagggtt 540
atcctggtcc tgtttcttt tgctggagtc gacgt 576

<210> 45

<211> 576

<212> DNA

<213> Homo sapiens

<220>

<223> Individual Isolate: SA1

<400> 45

gtcccttacc ggaatgcctc tgggtttac catgtcacca atgactgccc aaactcctcc 60
atagtctacg aggctgatag cctgatctt cacgcacccgt gctgcgtgcc ctgtgtcagg 120
caagataatg tcaagtaggtg ctgggtccaa atcaccccca cactgtcagc cccgaccttc 180
ggagcgggtca cggctctct tcggaggggcc gttgactact tagcgggagg agctgcttc 240

tgctccgcac tatacgtcgg cgacgcgtgc gggcagtgt ttctggtagg ccaaatgttc 300
acctataggc ctcgcagca taccacagtg caggactgca actgttccat ttacagtggc 360
catatcacccg gccaccggat ggcttggac atgatgatga attggtcacc tacgacagcc 420
ttgctgatgg cccagatgct acggatcccc caggtggtca tagacatcat agccggggc 480
cactgggggg tcttggtagc cgccgcatac ttgcgtcgg ccgccaactg ggctaaggta 540
tgctggtagc tgttcctgtt tgcgggggc gatggc 576

<210> 46
<211> 576
<212> DNA
<213> Homo sapiens

<220>
<223> Individual Isolate: SA4

<400> 46
gtccctacc gaaacgcctc tgggttat catgtcacca atgattgccaaactttcc 60
atagtttacg aggctgataa cctgatcttgcatgcacccgttgcgtgcc ttgtgtcagg 120
caagataatgc ttagtaagtgc tgggtccaa atcaccggccgttgcagcccaatctc 180
ggagcggtca cggctccctctc tggggggcc gttgactact tagcgggagg ggctgcctc 240
tgctccgcac tatacgtcgg ggacgcgtgc gggcagtgt ttggtagg ccaaatgttc 300
acctataggc ctcgcagca cactacggtgcagactgca attgctctat ttacagtggc 360
catatcacccg gccaccggat ggcattggac atgatgatga attggtcacc tacgacggcc 420
ttgctgatgg cccagttgtacggattccc caggtggtca tcgacatcat tgccggggc 480
cactgggggg tcttggtagc cgccgcatac ttgcgtcag ccgctaactg ggctaaggta 540
atactggtct tgtttctgtt tgcgggggc gatgcc 576

<210> 47
<211> 576
<212> DNA
<213> Homo sapiens

<220>
<223> Individual Isolate: SA5

<400> 47
gtccctacc gaaatgcctc tgggttat catgtcacca atgattgccaaactttcc 60
atagtttacg aggctgataa cctgattctgcacgcacccgttgcgtgcc ctgtgtcaag 120
gaaggtaatgc ttagtaggtgc tgggtccaa atcaccggccattgtcagcccaacctc 180
ggagcggtca cggctccctctc tggggggcc gttgactact tagcgggagg ggctgcctc 240
tgctccgcac tatacgtcgg ggacgcgtgc gggcagtgt ttggtagg ccaaatgttc 300
acctataggc ctcgcagca tactacggtgcagactgca actgttccat ttacagcggc 360
catatcacccg gccaccgaat ggcattggac atgatgatga attggtcacc tacgacagcc 420
ttggtgatgg cccaggtgtacggattccc caagtggtca ttgacatcat tgccggggc 480
cactgggggg tcttggtagc cgccgcatac ttgcgtcag ccgctaactg ggctaaggta 540
tgctggtagc tgtttctgtt tgcgggggc gatgcc 576

<210> 48
<211> 576
<212> DNA
<213> Homo sapiens

<400> 48
gttccttacc ggaatgcctc tgggtgtat catgttacca atgattgccaaactcttcc 60
atagtctatg aggctgatga cctgatccta cacgacacccgctg gctgcgtgcc ctgtgtccgg 120
aaggataatg tcagtagatg ctgggttcat atcaccggcca cactatcagc cccgagcctc 180
ggagcggtca cggctcctct tcggagggcc gttgattact tggcgggagg ggccgcctg 240
tgctccgcgt tatacgtcgg agacgtgtc gggcattgt ttttggtagg ccaaatgttc 300
acctataaggc ctcgcccagca tgctacggta caggactgca actgctccat ttacagtggc 360
catatcactg gccaccggat ggcacatggac atgatgtga attggtcacc cgcacagcc 420
ttgggtatgg cccaaatgtc acggattccc caggtggtca ttgacatcat tgccggggc 480
cactgggggg tcttggtcgc cgctgcatac ttgcgtcgg cggctaactg ggctaagggtt 540
gtgctggtct tgttctgtt tgccgggggtt gatgcc 576

<210> 49
<211> 576
<212> DNA
<213> Homo sapiens

<220>
<223> Individual Isolate: SA7

<400> 49
gtcccccttacc gaaatgcctc cggggtttat catgtcacca atgattgccaaactcttcc 60
atagtctatg aggctgacaa cctgatcctg cacgacacccgctg gttgcgtgcc ctgtgtcaga 120
caaataatg tcagtaggtg ctgggtccaa atcaccggcca cattgtcagc cccgaacccctc 180
ggagcggtca cggctcctct tcggagggcc gttgactacc tagcgggagg ggctgcctc 240
tgctccgcgc tatacgtcgg ggacgcgtc gggcagtgt ttttggtagg ccagatgttc 300
actataaggc ctcgcccagca cactacggta caggactgca actgctccat ttacagtggc 360
catatcacccg gccaccgaat ggcacatggac atgatgtga attggtcacc tacgacagcc 420
ttgggtatgg cccagttgtc acggattccc caggtggtca tcgacatcat tgccggggc 480
cactgggggg tcttggtcgc cgccgcatac ttgcgtcgg cggctaactg ggctaagggtt 540
gtgctggtct tgttctgtt tgccgggggtc gatgcc 576

<210> 50
<211> 576
<212> DNA
<213> Homo sapiens

<220>
<223> Individual Isolate: SA13

<400> 50
gttccctacc gaaatgcctc tgggtttat catgtcacca atgattgccca aactcttcc 60
atcgctacg aggctgatga cctgatctt cacgcacctg gttgcgtgcc ctgtgttagg 120
cagggtaatg tcagtaggtg ctgggtccag atcacccca cactgtcagc cccgagcctc 180
ggagcggtca cggctctct tcggagggcc gttgactact tagcgggggg ggctgccctt 240
tgctccgcgt tatacgtcgg agacgcgtgc gggcagtgt ttttgttagg tcaaatgttc 300
acctatagcc ctcgcggca taatgttgc caggactgca actgttccat ttacagtggc 360
cacatcacccg gccaccggat ggcatggac atgatgatga attggtcacc tacaacagct 420
tttgtatgg cccagttgtt acggattccc caggtggtca ttgacatcat tgccggggcc 480
cactgggggg tcttggtcgc cgccgcatac tacgcgtcgg cggctaactg ggccaagggtt 540
tgctggtcc tgttctgtt tgcgggggc gatgcc 576

<210> 51
<211> 576
<212> DNA
<213> Homo sapiens

<220>
<223> Individual Isolate: HK2

<400> 51
cttacctacg gcaactccag tggctatac catctcacaa atgattgccca aactccagc 60
atcgctgg aggccgatgc tatgatctt cattgcctg gatgcttgc ttgtgtgagg 120
gtcgatgatc ggtccacctg ttggcatgtt gtgacccca ccctggccat accaaatgct 180
tccacgcccc caacgggatt ccgcaggcat gtggatctt ttgcgggccc cgcaagtgggtt 240
tgctcatccc tgtacatcg ggacctgtt ggctctctt tttggccggg acaactattc 300
accttcagc cccgcgtca ttggactgtt caagactgca actgctccat ctatacaggc 360
cacgtcaccc gccacaggat ggcttggac atgatgatga actggtcacc cacaaccact 420
ctggctctat ctagcatctt gagggatccct gagatttgtt cgagtgtat atttgtggc 480
cattggggga tactactagc cgttgcctac tttggcatgg ctggcaactg gctaaaagtt 540
ctggctgttc tgttcctatt tgcgagggtt gaagca 576

<210> 52
<211> 192
<212> PRT
<213> Homo sapiens

<220>
<223> Individual Isolate: DK7

<400> 52
Tyr Gln Val Arg Asn Ser Thr Gly Leu Tyr His Val Thr Asn Asp Cys
1 5 10 15

Pro Asn Ser Ser Ile Val Tyr Glu Ala Ala Asp Ala Ile Leu His Thr

20

25

30

Pro Gly Cys Val Pro Cys Val Arg Glu Gly Asn Val Ser Arg Cys Trp
 35 40 45

Val Ala Met Thr Pro Thr Val Ala Thr Arg Asp Gly Lys Leu Pro Thr
 50 55 60

Ala Gln Leu Arg Arg His Ile Asp Leu Leu Val Gly Ser Ala Thr Leu
 65 70 75 80

Cys Ser Ala Leu Tyr Val Gly Asp Leu Cys Gly Ser Val Phe Leu Val
 85 90 95

Gly Gln Leu Phe Thr Phe Ser Pro Arg Arg His Trp Thr Thr Gln Gly
 100 105 110

Cys Asn Cys Ser Ile Tyr Pro Gly His Ile Thr Gly His Arg Met Ala
 115 120 125

Trp Asp Met Met Met Asn Trp Ser Pro Thr Thr Ala Leu Val Val Ala
 130 135 140

Gln Leu Leu Arg Ile Pro Gln Ala Ile Leu Asp Met Ile Ala Gly Ala
 145 150 155 160

His Trp Gly Val Leu Ala Gly Ile Ala Tyr Phe Ser Met Val Gly Asn
 165 170 175

Trp Ala Lys Val Leu Val Val Leu Leu Phe Ala Gly Val Asp Ala
 180 185 190

<210> 53

<211> 192

<212> PRT

<213> Homo sapiens

<220>

<223> Individual Isolate: DK9

<400> 53

Tyr Gln Val Arg Asn Ser Ser Gly Leu Tyr His Val Thr Asn Asp Cys
 1 5 10 15

Pro	Asn	Ser	Ser	Ile	Val	Tyr	Glu	Ala	Ala	Asp	Ala	Ile	Leu	His	Ser
20							25							30	
Pro	Gly	Cys	Val	Pro	Cys	Val	Arg	Glu	Gly	Asn	Ala	Ser	Lys	Cys	Trp
35							40							45	
Val	Ala	Val	Ala	Pro	Thr	Val	Ala	Thr	Arg	Asp	Gly	Lys	Leu	Pro	Ala
50							55						60		
Thr	Gln	Leu	Arg	Arg	His	Ile	Asp	Leu	Leu	Val	Gly	Ser	Ala	Thr	Leu
65							70						75		80
Cys	Ser	Ala	Leu	Tyr	Val	Gly	Asp	Leu	Cys	Gly	Ser	Val	Phe	Leu	Val
85							90						95		
Gly	Gln	Leu	Phe	Thr	Phe	Ser	Pro	Arg	Arg	His	Trp	Thr	Thr	Gln	Asp
100							105						110		
Cys	Asn	Cys	Ser	Ile	Tyr	Pro	Gly	His	Ile	Thr	Gly	His	Arg	Met	Ala
115							120						125		
Trp	Asp	Met	Met	Met	Asn	Trp	Ser	Pro	Thr	Ala	Ala	Leu	Val	Met	Ala
130							135						140		
Gln	Leu	Leu	Arg	Ile	Pro	Gln	Ala	Ile	Leu	Asp	Met	Ile	Ala	Gly	Ala
145							150						155		160
His	Trp	Gly	Val	Leu	Ala	Gly	Ile	Ala	Tyr	Phe	Ser	Met	Val	Gly	Asn
165							170						175		
Trp	Ala	Lys	Val	Val	Val	Leu	Leu	Leu	Phe	Thr	Gly	Val	Asp	Ala	
180							185						190		

<210> 54
 <211> 192
 <212> PRT
 <213> Homo sapiens

<220>
 <223> Individual Isolate: DR1

<400> 54
 His Gln Val Arg Asn Ser Thr Gly Leu Tyr His Val Thr Asn Asp Cys
 1 5 10 15

Pro Asn Ser Ser Ile Val Tyr Glu Ala Ala Asp Ala Ile Leu His Ala		
20	25	30
Pro Gly Cys Val Pro Cys Val Arg Glu Gly Asn Ala Ser Arg Cys Trp		
35	40	45
Val Ala Val Thr Pro Thr Val Ala Thr Arg Asp Gly Lys Leu Pro Thr		
50	55	60
Thr Gln Leu Arg Arg His Ile Asp Leu Leu Val Gly Ser Ala Thr Leu		
65	70	75
80		
Cys Ser Ala Leu Tyr Val Gly Asp Leu Cys Gly Ser Val Phe Leu Val		
85	90	95
Gly Gln Leu Phe Thr Phe Ser Pro Arg Arg His Trp Thr Thr Gln Asp		
100	105	110
Cys Asn Cys Ser Ile Tyr Pro Gly His Ile Thr Gly His Arg Met Ala		
115	120	125
Trp Asp Met Met Asn Trp Ser Pro Thr Thr Ala Leu Val Met Ala		
130	135	140
Gln Leu Leu Arg Ile Pro Gln Ala Ile Leu Asp Met Ile Ala Gly Ala		
145	150	155
160		
His Trp Gly Val Leu Ala Gly Ile Ala Tyr Phe Ser Met Val Gly Asn		
165	170	175
Trp Ala Lys Val Val Val Leu Leu Leu Phe Ala Gly Val Asp Ala		
180	185	190

<210> 55
 <211> 192
 <212> PRT
 <213> Homo sapiens

<220>
 <223> Individual Isolate: DR4

<400> 55
 His Gln Val Arg Asn Ser Thr Gly Leu Tyr His Val Thr Asn Asp Cys

1	5	10	15
Pro Asn Ser Ser Ile Val Tyr Glu Ala Ala Asp Ala Ile Leu His Thr			
20	25	30	
Pro Gly Cys Val Pro Cys Val Arg Glu Gly Asn Thr Ser Arg Cys Trp			
35	40	45	
Val Ala Val Thr Pro Thr Val Ala Thr Arg Asp Gly Lys Leu Pro Thr			
50	55	60	
Thr Gln Leu Arg Arg His Ile Asp Leu Leu Val Gly Ser Ala Thr Leu			
65	70	75	80
Cys Ser Ala Leu Tyr Val Gly Asp Leu Cys Gly Ser Val Phe Leu Val			
85	90	95	
Gly Gln Leu Phe Thr Phe Ser Pro Arg His His Trp Thr Thr Gln Asp			
100	105	110	
Cys Asn Cys Ser Ile Tyr Pro Gly His Ile Thr Gly His Arg Met Ala			
115	120	125	
Trp Asp Met Met Met Asn Trp Ser Pro Thr Thr Ala Leu Val Val Ala			
130	135	140	
Gln Leu Leu Arg Ile Pro Gln Ala Ile Leu Asp Met Ile Ala Gly Ala			
145	150	155	160
His Trp Gly Val Leu Ala Gly Ile Ala Tyr Phe Ser Met Val Gly Asn			
165	170	175	
Trp Ala Lys Val Leu Val Val Leu Leu Leu Phe Ala Gly Val Asp Ala			
180	185	190	

<210> 56
 <211> 192
 <212> PRT
 <213> Homo sapiens

 <220>
 <223> Individual Isolate: S14

 <400> 56

Tyr	Gln	Val	Arg	Asn	Ser	Thr	Gly	Leu	Tyr	His	Val	Thr	Asn	Asp	Cys
1		5				10						15			
Pro	Asn	Ser	Ser	Ile	Val	Tyr	Glu	Thr	Ala	Asp	Ala	Ile	Leu	His	Ala
	20					25						30			
Pro	Gly	Cys	Val	Pro	Cys	Val	Arg	Glu	Gly	Asn	Thr	Ser	Arg	Cys	Trp
	35					40						45			
Val	Ala	Met	Thr	Pro	Thr	Val	Ala	Thr	Arg	Asp	Gly	Lys	Leu	Pro	Ala
	50					55						60			
Thr	Gln	Leu	Arg	Arg	Tyr	Ile	Asp	Leu	Leu	Val	Gly	Ser	Ala	Thr	Leu
	65				70					75				80	
Cys	Ser	Ala	Leu	Tyr	Val	Gly	Asp	Leu	Cys	Gly	Ser	Val	Phe	Leu	Val
	85					90						95			
Gly	Gln	Leu	Phe	Thr	Phe	Ser	Pro	Arg	Arg	Leu	Trp	Thr	Thr	Gln	Asp
	100				105							110			
Cys	Asn	Cys	Ser	Ile	Tyr	Pro	Gly	His	Ile	Thr	Gly	His	Arg	Met	Ala
	115					120					125				
Trp	Asp	Met	Met	Met	Asn	Trp	Ser	Pro	Thr	Thr	Ala	Leu	Val	Val	Ala
	130					135					140				
Gln	Leu	Leu	Arg	Ile	Pro	Gln	Ala	Ile	Leu	Asp	Met	Ile	Ala	Gly	Ala
	145				150					155			160		
His	Trp	Gly	Val	Leu	Ala	Gly	Ile	Ala	Tyr	Phe	Ser	Met	Val	Gly	Asn
	165					170						175			
Trp	Ala	Lys	Val	Leu	Val	Val	Leu	Leu	Leu	Phe	Ala	Gly	Val	Asp	Ala
	180					185						190			

<210> 57
 <211> 192
 <212> PRT
 <213> Homo sapiens

<220>
 <223> Individual Isolate: S18

<400> 57

Tyr Gln Val Arg Asn Ser Thr Gly Leu Tyr His Val Thr Asn Asp Cys
1 5 10 15

Pro Asn Ser Ser Ile Val Tyr Glu Thr Ala Asp Thr Ile Leu His Ser
20 25 30

Pro Gly Cys Val Pro Cys Val Arg Glu Gly Asn Ala Ser Arg Cys Trp
35 40 45

Val Pro Val Ala Pro Thr Val Ala Thr Arg Asp Gly Lys Leu Pro Ala
50 55 60

Thr Gln Leu Arg Arg His Ile Asp Leu Leu Val Gly Ser Ala Thr Leu
65 70 75 80

Cys Ser Ala Leu Tyr Val Gly Asp Leu Cys Gly Ser Val Phe Leu Val
85 90 95

Ser Gln Leu Phe Thr Ile Ser Pro Arg Arg His Trp Thr Thr Gln Asp
100 105 110

Cys Asn Cys Ser Ile Tyr Pro Gly His Ile Thr Gly His Arg Met Ala
115 120 125

Trp Asp Met Met Met Asn Trp Ser Pro Thr Thr Ala Leu Val Ile Ala
130 135 140

Gln Leu Leu Arg Val Pro Gln Ala Val Leu Asp Met Ile Ala Gly Ala
145 150 155 160

His Trp Gly Val Leu Ala Gly Ile Ala Tyr Phe Ser Met Ala Gly Asn
165 170 175

Trp Ala Lys Val Leu Leu Val Leu Leu Phe Ala Gly Val Asp Ala
180 185 190

<210> 58

<211> 192

<212> PRT

<213> Homo sapiens

<220>

<223> Individual Isolate: SW1

<400> 58

Tyr Gln Val Arg Asn Ser Ser Gly Leu Tyr His Val Thr Asn Asp Cys
1 5 10 15

Pro Asn Ser Ser Ile Val Tyr Glu Thr Ala Asp Ala Ile Leu His Ser
20 25 30

Pro Gly Cys Val Pro Cys Val Arg Glu Asp Gly Ala Pro Lys Cys Trp
35 40 45

Val Ala Val Ala Pro Thr Val Ala Thr Arg Asp Gly Lys Leu Pro Ala
50 55 60

Thr Gln Leu Arg Arg His Ile Asp Leu Leu Val Gly Ser Ala Thr Leu
65 70 75 80

Cys Ser Ala Leu Tyr Val Gly Asp Leu Cys Gly Ser Val Phe Leu Val
85 90 95

Ser Gln Leu Phe Thr Phe Ser Pro Arg Arg His Trp Thr Thr Gln Asp
100 105 110

Cys Asn Cys Ser Ile Tyr Pro Gly His Ile Thr Gly His Arg Met Ala
115 120 125

Trp Asp Met Met Asn Trp Ser Pro Thr Thr Ala Leu Val Val Ala
130 135 140

Gln Leu Leu Arg Ile Pro Gln Ala Val Leu Asp Met Ile Ala Gly Ala
145 150 155 160

His Trp Gly Val Leu Ala Gly Ile Ala Tyr Phe Ser Met Val Gly Asn
165 170 175

Trp Ala Lys Val Leu Ile Val Leu Leu Phe Ser Gly Val Asp Ala
180 185 190

<210> 59

<211> 192

<212> PRT

<213> Homo sapiens

<220>

<223> Individual Isolate: US11

<400> 59

Tyr Gln Val Arg Asn Ser Thr Gly Leu Tyr His Val Thr Asn Asp Cys
1 5 10 15

Pro Asn Ser Ser Ile Val Tyr Glu Ala Ala Asp Ala Ile Leu His Thr
20 25 30

Pro Gly Cys Val Pro Cys Val Arg Glu Gly Asn Ala Ser Arg Cys Trp
35 40 45

Val Ala Met Thr Pro Thr Val Ala Thr Arg Asp Gly Lys Leu Pro Thr
50 55 60

Thr Gln Leu Arg Arg His Ile Asp Leu Leu Val Gly Ser Ala Thr Leu
65 70 75 80

Cys Ser Ala Leu Tyr Val Gly Asp Leu Cys Gly Ser Val Phe Leu Val
85 90 95

Gly Gln Leu Phe Thr Phe Ser Pro Arg Arg His Trp Thr Thr Gln Gly
100 105 110

Cys Asn Cys Ser Ile Tyr Pro Gly His Ile Thr Gly His Arg Met Ala
115 120 125

Trp Asp Met Met Asn Trp Ser Pro Thr Ala Ala Leu Val Val Ala
130 135 140

Gln Leu Leu Arg Ile Pro Gln Ala Ile Leu Asp Met Ile Ala Gly Ala
145 150 155 160

His Trp Gly Val Leu Ala Gly Ile Ala Tyr Phe Ser Met Val Gly Asn
165 170 175

Trp Ala Lys Val Leu Val Val Leu Leu Phe Ala Gly Val Asp Ala
180 185 190

<210> 60

<211> 192

<212> PRT

<213> Homo sapiens

<220>

<223> Individual Isolate: D1

<400> 60

Tyr Glu Val Arg Asn Val Ser Gly Val Tyr His Val Thr Asn Asp Cys
1 5 10 15

Ser Asn Ser Ser Ile Val Tyr Glu Thr Ala Asp Met Ile Met His Thr
20 25 30

Pro Gly Cys Val Pro Cys Val Arg Glu Asp Asn Ser Ser Arg Cys Trp
35 40 45

Val Ala Leu Thr Pro Thr Leu Ala Ala Arg Asn Gly Asn Val Pro Thr
50 55 60

Thr Ala Ile Arg Arg His Val Asp Leu Leu Val Gly Ala Ala Ala Phe
65 70 75 80

Cys Ser Ala Met Tyr Val Gly Asp Leu Cys Gly Ser Val Phe Leu Ile
85 90 95

Ser Gln Leu Phe Thr Leu Ser Pro Arg Arg His Glu Thr Val Gln Glu
100 105 110

Cys Asn Cys Ser Ile Tyr Pro Gly His Val Thr Gly His Arg Met Ala
115 120 125

Trp Asp Met Met Met Asn Trp Ser Pro Thr Thr Ala Leu Val Val Ser
130 135 140

Gln Leu Leu Arg Ile Pro Gln Ala Val Met Asp Met Val Ala Gly Ala
145 150 155 160

His Trp Gly Val Leu Ala Gly Leu Ala Tyr Tyr Ser Met Val Gly Asn
165 170 175

Trp Ala Lys Val Leu Ile Val Met Leu Leu Phe Ala Gly Val Asp Gly
180 185 190

<210> 61

<211> 192

<212> PRT

<213> Homo sapiens

<220>

<223> Individual Isolate: D3

<400> 61

Tyr Glu Val Arg Asn Val Ser Gly Val Tyr Gln Val Thr Asn Asp Cys
1 5 10 15

Ser Asn Ser Ser Ile Val Tyr Glu Thr Ala Asp Met Ile Met His Thr
20 25 30

Pro Gly Cys Val Pro Cys Val Arg Glu Asp Asn Ser Ser Arg Cys Trp
35 40 45

Val Ala Leu Thr Pro Thr Leu Ala Ala Arg Asn Ser Ser Val Pro Thr
50 55 60

Thr Thr Ile Arg Arg His Val Asp Leu Leu Val Gly Ala Ala Ala Phe
65 70 75 80

Cys Ser Ala Met Tyr Val Gly Asp Leu Cys Gly Ser Val Phe Leu Val
85 90 95

Ser Gln Leu Phe Thr Phe Ser Pro Arg Arg His Glu Thr Val Gln Glu
100 105 110

Cys Asn Cys Ser Ile Tyr Pro Gly His Val Thr Gly His Arg Met Ala
115 120 125

Trp Asp Met Met Asn Trp Ser Pro Thr Ala Ala Leu Val Val Ser
130 135 140

Gln Leu Leu Arg Ile Pro Gln Ala Val Val Asp Met Val Ala Gly Ala
145 150 155 160

His Trp Gly Val Leu Ala Gly Leu Ala Tyr Tyr Ser Met Val Gly Asn
165 170 175

Trp Ala Lys Val Leu Ile Val Met Leu Leu Phe Ala Gly Val Asp Gly
180 185 190

<210> 62

<211> 192

<212> PRT

<213> Homo sapiens

<220>

<223> Individual Isolate: DK1

<400> 62

Tyr Glu Val Arg Asn Val Ser Gly Val Tyr His Val Thr Asn Asp Cys
1 5 10 15

Ser Asn Ser Ser Ile Val Tyr Glu Ala Val Asp Val Ile Met His Thr
20 25 30

Pro Gly Cys Val Pro Cys Val Arg Glu Asn Asn His Ser Arg Cys Trp
35 40 45

Val Ala Leu Thr Pro Thr Leu Ala Ala Arg Asn Ala Ser Ile Pro Thr
50 55 60

Thr Thr Ile Arg Arg His Val Asp Leu Leu Val Gly Ala Ala Ala Phe
65 70 75 80

Cys Ser Ala Met Tyr Val Gly Asp Leu Cys Gly Ser Val Phe Leu Val
85 90 95

Ser Gln Leu Phe Thr Phe Ser Pro Arg Arg His Glu Thr Ala Gln Asp
100 105 110

Cys Asn Cys Ser Ile Tyr Pro Gly His Val Ser Gly His Arg Met Ala
115 120 125

Trp Asp Met Met Asn Trp Ser Pro Thr Thr Ala Leu Val Leu Ser
130 135 140

Gln Leu Leu Arg Ile Pro Gln Ala Val Val Asp Met Val Ala Gly Ala
145 150 155 160

His Trp Gly Val Leu Ala Gly Leu Ala Tyr Tyr Ser Met Ala Gly Asn
165 170 175

Trp Ala Lys Val Leu Ile Val Leu Leu Phe Ala Gly Val Asp Gly
180 185 190

<210> 63

<211> 192

<212> PRT

<213> Homo sapiens

<220>

<223> Individual Isolate: HK3

<400> 63

Tyr Glu Val Arg Asn Val Ser Gly Ile Tyr His Val Thr Asn Asp Cys
1 5 10 15

Ser Asn Ser Ser Val Val Tyr Glu Thr Ala Asp Met Ile Met His Thr
20 25 30

Pro Gly Cys Val Pro Cys Val Arg Glu Asn Asn Ser Ser Arg Cys Trp
35 40 45

Val Ala Leu Thr Pro Thr Leu Ala Ala Arg Asn Val Ser Val Pro Thr
50 55 60

Thr Thr Ile Arg Arg His Val Asp Leu Leu Val Gly Ala Ala Ala Phe
65 70 75 80

Cys Ser Ala Met Tyr Val Gly Asp Leu Cys Gly Ser Val Phe Leu Val
85 90 95

Ser Gln Leu Phe Thr Phe Ser Pro Arg Arg His Glu Thr Val Gln Asp
100 105 110

Cys Asn Cys Ser Leu Tyr Pro Gly His Val Ser Gly His Arg Met Ala
115 120 125

Trp Asp Met Met Met Asn Trp Ser Pro Thr Ala Ala Leu Val Val Ser
130 135 140

Gln Leu Leu Arg Ile Pro Gln Ala Val Val Asp Met Val Ala Gly Ala
145 150 155 160

His Trp Gly Val Leu Ala Gly Leu Ala Tyr Tyr Ser Met Val Gly Asn
165 170 175

Trp Ala Lys Val Leu Ile Val Met Leu Leu Phe Ala Gly Val Asp Gly
180 185 190

<210> 64

<211> 192
<212> PRT
<213> Homo sapiens

<220>
<223> Individual Isolate: HK4

<400> 64

His Glu Val His Asn Val Ser Gly Ile Tyr His Val Thr Asn Asp Cys
1 5 10 15

Ser Asn Ser Ser Ile Val Tyr Glu Ala Ala Asp Met Ile Met His Thr
20 25 30

Pro Gly Cys Val Pro Cys Val Arg Glu Asn Asn Ser Ser Arg Cys Trp
35 40 45

Val Ala Leu Thr Pro Thr Leu Ala Ala Arg Asn Ala Ser Ile Pro Thr
50 55 60

Thr Thr Ile Arg Arg His Val Asp Leu Leu Val Gly Ala Ala Ala Phe
65 70 75 80

Cys Ser Ala Met Tyr Val Gly Asp Leu Cys Gly Ser Val Phe Leu Val
85 90 95

Ser Gln Leu Phe Thr Phe Ser Pro Arg Arg His Glu Thr Val Gln Asp
100 105 110

Cys Asn Cys Ser Ile Tyr Pro Gly His Val Ser Gly His Arg Met Ala
115 120 125

Trp Asp Met Met Asn Trp Ser Pro Thr Ala Ala Leu Val Val Ser
130 135 140

Gln Leu Leu Arg Leu Pro Gln Ala Val Met Asp Met Val Ala Gly Ala
145 150 155 160

His Trp Gly Val Leu Ala Gly Leu Ala Tyr Tyr Ser Met Val Gly Asn
165 170 175

Trp Ala Lys Val Leu Ile Val Met Leu Leu Phe Ala Gly Val Asp Gly
180 185 190

<210> 65
<211> 192
<212> PRT
<213> Homo sapiens

<220>
<223> Individual Isolate: HK5

<400> 65
Tyr Glu Val Arg Asn Val Ser Gly Val Tyr His Val Thr Asn Asp Cys
1 5 10 15

Ser Asn Leu Ser Ile Val Tyr Glu Thr Thr Asp Met Ile Met His Thr
20 25 30

Pro Gly Cys Val Pro Cys Val Arg Glu Asn Asn Ser Ser Arg Cys Trp
35 40 45

Val Ala Leu Ala Pro Thr Leu Ala Ala Arg Asn Ala Ser Val Pro Thr
50 55 60

Thr Ala Ile Arg Arg His Val Asp Leu Leu Val Gly Ala Ala Ala Phe
65 70 75 80

Cys Ser Ala Met Tyr Val Gly Asp Leu Cys Gly Ser Val Phe Leu Val
85 90 95

Ser Gln Leu Phe Thr Phe Ser Pro Arg Arg His Glu Thr Val Gln Asp
100 105 110

Cys Asn Cys Ser Ile Tyr Pro Gly His Val Thr Gly His Arg Met Ala
115 120 125

Trp Asp Met Met Met Asn Trp Ser Pro Thr Thr Ala Leu Val Val Ser
130 135 140

Gln Leu Leu Arg Ile Pro Gln Ala Val Val Asp Met Val Ala Gly Ala
145 150 155 160

His Trp Gly Val Leu Ala Gly Leu Ala Tyr Tyr Ser Met Val Gly Asn
165 170 175

Trp Ala Lys Val Leu Ile Val Met Leu Leu Phe Ala Gly Val Asp Gly
180 185 190

<210> 66
<211> 192
<212> PRT
<213> Homo sapiens

<220>
<223> Individual Isolate: HK8

<400> 66
Tyr Glu Val Arg Asn Val Ser Gly Ile Tyr His Val Thr Asn Asp Cys
1 5 10 15

Ser Asn Ser Ser Ile Val Tyr Glu Thr Ala Asp Met Ile Met His Thr
20 25 30

Pro Gly Cys Met Pro Cys Val Arg Glu Asn Asn Ser Ser Arg Cys Trp
35 40 45

Val Ala Leu Thr Pro Thr Leu Ala Ala Arg Asn Val Ser Val Pro Thr
50 55 60

Thr Thr Ile Arg Arg His Val Asp Leu Leu Val Gly Ala Ala Ala Phe
65 70 75 80

Cys Ser Ala Met Tyr Val Gly Asp Leu Cys Gly Ser Val Phe Leu Val
85 90 95

Ser Gln Leu Phe Thr Phe Ser Pro Arg Arg His Glu Thr Val Gln Asp
100 105 110

Cys Asn Cys Ser Ile Tyr Pro Gly His Val Ser Gly His Arg Met Ala
115 120 125

Trp Asp Met Met Met Asn Trp Ser Pro Thr Thr Ala Leu Val Val Ser
130 135 140

Gln Leu Leu Arg Ile Pro Gln Ala Ile Val Asp Met Val Ala Gly Ala
145 150 155 160

His Trp Gly Val Leu Ala Gly Leu Ala Tyr Tyr Ser Met Val Gly Asn
165 170 175

Trp Ala Lys Val Leu Ile Val Met Leu Leu Phe Ala Gly Val Asp Gly
180 185 190

<210> 67
<211> 192
<212> PRT
<213> Homo sapiens

<220>
<223> Individual Isolate: IND5

<400> 67
Tyr Glu Val Arg Asn Val Ser Gly Val Tyr His Val Thr Asn Asp Cys
1 5 10 15

Ser Asn Ser Ser Ile Val Tyr Glu Ala Ala Asp Met Ile Met His Thr
20 25 30

Pro Gly Cys Val Pro Cys Val Arg Glu Gly Asn Ser Ser Arg Cys Trp
35 40 45

Val Ala Leu Thr Pro Thr Leu Ala Ala Arg Asn Ala Ser Val Ser Thr
50 55 60

Thr Thr Ile Arg His His Val Asp Leu Leu Val Gly Ala Ala Ala Phe
65 70 75 80

Cys Ser Ala Met Tyr Val Gly Asp Leu Cys Gly Ser Val Phe Leu Val
85 90 95

Ser Gln Leu Phe Thr Phe Ser Pro Arg Arg His Glu Thr Val Gln Asp
100 105 110

Cys Asn Cys Ser Ile Tyr Pro Gly His Val Ser Gly His Arg Met Ala
115 120 125

Trp Asp Met Met Met Asn Trp Ser Pro Thr Ala Ala Leu Val Val Ser
130 135 140

Gln Leu Leu Arg Ile Pro Gln Ala Val Val Asp Met Val Ala Gly Ala
145 150 155 160

His Trp Gly Ile Leu Ala Gly Leu Ala Tyr Tyr Ser Met Val Gly Asn
165 170 175

Trp Ala Lys Val Leu Ile Val Met Leu Leu Phe Ala Gly Val Asp Gly
180 185 190

<210> 68
<211> 192
<212> PRT
<213> Homo sapiens

<220>
<223> Individual Isolate: IND8

<400> 68
Tyr Glu Val Arg Asn Val Ser Gly Val Tyr His Val Thr Asn Asp Cys
1 5 10 15
Ser Asn Ser Ser Ile Val Tyr Glu Ala Ala Asp Met Ile Met His Thr
20 25 30
Pro Gly Cys Val Pro Cys Val Arg Glu Gly Asn Phe Ser Ser Cys Trp
35 40 45
Val Ala Leu Thr Pro Thr Leu Ala Ala Arg Asn Ala Ser Val Pro Thr
50 55 60
Thr Thr Ile Arg Arg His Val Asp Leu Leu Val Gly Ala Ala Ala Phe
65 70 75 80
Cys Ser Ala Met Tyr Val Gly Asp Leu Cys Gly Ser Val Phe Leu Val
85 90 95
Ser Gln Leu Phe Thr Phe Ser Pro Arg Arg His Glu Thr Val Gln Asp
100 105 110
Cys Asn Cys Ser Ile Tyr Pro Gly His Val Ser Gly His Arg Met Ala
115 120 125
Trp Asp Met Met Met Asn Trp Ser Pro Thr Ala Ala Leu Val Val Ser
130 135 140
Gln Leu Leu Arg Ile Pro Gln Ala Val Val Asp Met Val Ala Gly Ala
145 150 155 160
His Trp Gly Ile Leu Ala Gly Leu Ala Tyr Tyr Ser Met Val Gly Asn
165 170 175
Trp Ala Lys Val Leu Ile Val Met Leu Leu Phe Ala Gly Val Asp Gly
180 185 190

<210> 69
<211> 192
<212> PRT
<213> Homo sapiens

<220>
<223> Individual Isolate: P10

<400> 69
Tyr Glu Val Arg Asn Val Ser Gly Val Tyr His Val Thr Asn Asp Cys
1 5 10 15

Ser Asn Ser Ser Ile Val Tyr Glu Ala Ala Asp Met Ile Met His Thr
20 25 30

Pro Gly Cys Val Pro Cys Val Arg Glu Asn Asn Ser Ser Arg Cys Trp
35 40 45

Val Ala Leu Thr Pro Thr Leu Ala Ala Arg Asn Ser Ser Val Pro Thr
50 55 60

Thr Ala Ile Arg Arg His Val Asp Leu Leu Val Gly Ala Ala Ala Phe
65 70 75 80

Cys Ser Ala Met Tyr Val Gly Asp Leu Cys Gly Ser Val Leu Leu Val
85 90 95

Ser Gln Leu Phe Thr Phe Ser Pro Arg Arg His Trp Thr Val Gln Asp
100 105 110

Cys Asn Cys Ser Ile Tyr Pro Gly His Val Ser Gly His Arg Met Ala
115 120 125

Trp Asp Met Met Met Asn Trp Ser Pro Thr Ala Ala Leu Val Val Ser
130 135 140

Gln Leu Leu Arg Ile Pro Gln Ala Ile Leu Asp Val Val Ala Gly Ala
145 150 155 160

His Trp Gly Val Leu Ala Gly Leu Ala Tyr Tyr Ser Met Val Gly Asn
165 170 175

Trp Ala Lys Val Leu Ile Val Met Leu Leu Phe Ala Gly Val Asp Gly
180 185 190

<210> 70
<211> 192
<212> PRT
<213> Homo sapiens

<220>
<223> Individual Isolate: S9

<400> 70
Tyr Glu Val Arg Asn Val Ser Gly Ala Tyr His Val Thr Asn Asp Cys
1 5 10 15

Ser Asn Ser Ser Ile Val Tyr Glu Ala Ala Asp Val Ile Met His Thr
20 25 30

Pro Gly Cys Val Pro Cys Val Gln Glu Gly Asn Ser Ser Gln Cys Trp
35 40 45

Val Ala Leu Thr Pro Thr Leu Ala Ala Arg Asn Ala Thr Val Pro Thr
50 55 60

Thr Thr Ile Arg Arg His Val Asp Leu Leu Val Gly Ala Ala Val Phe
65 70 75 80

Cys Ser Ala Met Tyr Val Gly Asp Leu Cys Gly Ser Val Phe Leu Ile
85 90 95

Ser Gln Leu Phe Thr Ile Ser Pro Arg Arg His Glu Thr Val Gln Asn
100 105 110

Cys Asn Cys Ser Ile Tyr Pro Gly His Val Thr Gly His Arg Met Ala
115 120 125

Trp Asp Met Met Met Asn Trp Ser Pro Thr Thr Ala Leu Val Val Ser
130 135 140

Gln Leu Leu Arg Ile Pro Gln Ala Val Met Asp Met Val Ala Gly Ala
145 150 155 160

His Trp Gly Val Leu Ala Gly Leu Ala Tyr Tyr Ser Met Val Gly Asn
165 170 175

Trp Ala Lys Val Leu Ile Val Met Leu Leu Phe Ala Gly Val Asp Gly

180

185

190

<210> 71
<211> 192
<212> PRT
<213> Homo sapiens

<220>
<223> Individual Isolate: S45

<400> 71
Tyr Glu Val Arg Asn Val Ser Gly Ala Tyr His Val Thr Asn Asp Cys
1 5 10 15

Ser Asn Ser Ser Ile Val Tyr Glu Ala Val Asp Val Ile Leu His Thr
20 25 30

Pro Gly Cys Val Pro Cys Val Arg Glu Asn Asn Ser Ser Arg Cys Trp
35 40 45

Val Ala Leu Thr Pro Thr Leu Ala Ala Arg Asn Ser Ser Val Pro Thr
50 55 60

Thr Thr Ile Arg Arg His Val Asp Leu Leu Val Gly Ala Ala Ala Phe
65 70 75 80

Cys Ser Ala Met Tyr Val Gly Asp Leu Cys Gly Ser Val Phe Leu Val
85 90 95

Ser Gln Leu Phe Thr Phe Ser Pro Arg Arg His Glu Thr Val Gln Asp
100 105 110

Cys Asn Cys Ser Ile Tyr Pro Gly His Val Thr Gly His Arg Met Ala
115 120 125

Trp Asp Met Met Met Asn Trp Ser Pro Thr Ala Ala Leu Val Val Ser
130 135 140

Gln Leu Leu Arg Ile Pro Gln Ala Val Val Asp Met Val Ala Gly Ala
145 150 155 160

His Trp Gly Val Leu Ala Gly Leu Ala Tyr Tyr Ser Met Val Gly Asn
165 170 175

Trp Ala Lys Val Leu Ile Val Met Leu Leu Phe Ala Gly Val Asp Gly
180 185 190

<210> 72

<211> 192

<212> PRT

<213> Homo sapiens

<220>

<223> Individual Isolate: SA10

<400> 72

Tyr Glu Val Arg Asn Val Ser Gly Met Tyr His Val Thr Asn Asp Cys
1 5 10 15

Ser Asn Ser Ser Ile Val Tyr Glu Ala Ala Asp Met Ile Met His Thr
20 25 30

Pro Gly Cys Val Pro Cys Val Arg Glu Asn Asn Ser Ser Arg Cys Trp
35 40 45

Val Ala Leu Thr Pro Thr Leu Ala Ala Arg Asn Ser Ser Val Pro Thr
50 55 60

Thr Thr Ile Arg Arg His Val Asp Leu Leu Val Gly Ala Ala Ala Phe
65 70 75 80

Cys Ser Ala Met Tyr Val Gly Asp Leu Cys Gly Ser Val Phe Leu Val
85 90 95

Ser Gln Leu Phe Thr Phe Ser Pro Arg Arg Tyr Glu Thr Val Gln Asp
100 105 110

Cys Asn Cys Ser Ile Tyr Pro Gly Arg Val Thr Gly His Arg Met Ala
115 120 125

Trp Asp Met Met Asn Trp Ser Pro Thr Thr Ala Leu Val Val Ser
130 135 140

Gln Leu Leu Arg Ile Pro Gln Ala Ile Val Asp Met Val Ala Gly Ala
145 150 155 160

His Trp Gly Val Leu Ala Gly Leu Ala Tyr Tyr Ser Met Val Gly Asn
165 170 175

Trp Ala Lys Val Leu Ile Val Met Leu Leu Phe Ala Gly Val Asp Gly
180 185 190

<210> 73
<211> 192
<212> PRT
<213> Homo sapiens

<220>
<223> Individual Isolate: SW2

<400> 73
Tyr Glu Val Arg Asn Val Ser Gly Val Tyr His Val Thr Asn Asp Cys
1 5 10 15

Ser Asn Ser Ser Ile Val Tyr Glu Thr Ala Asp Met Ile Met His Thr
20 25 30

Pro Gly Cys Val Pro Cys Val Arg Glu Ala Asn Ser Ser Arg Cys Trp
35 40 45

Val Ala Leu Thr Pro Thr Leu Ala Ala Arg Asn Thr Ser Val Pro Thr
50 55 60

Thr Thr Ile Arg Arg His Val Asp Leu Leu Val Gly Ala Ala Ala Phe
65 70 75 80

Cys Ser Val Met Tyr Val Gly Asp Leu Cys Gly Ser Val Phe Leu Val
85 90 95

Ser Gln Leu Phe Thr Phe Ser Pro Arg Arg His Glu Thr Val Gln Asp
100 105 110

Cys Asn Cys Ser Ile Tyr Pro Gly His Val Ser Gly His Arg Met Ala
115 120 125

Trp Asp Met Met Asn Trp Ser Pro Thr Ala Ala Leu Val Val Ser
130 135 140

Gln Leu Leu Arg Ile Pro Gln Ala Val Val Asp Met Val Ala Gly Ala
145 150 155 160

His Trp Gly Val Leu Ala Gly Leu Ala Tyr Tyr Ser Met Val Gly Asn

165

170

175

Trp Ala Lys Val Leu Ile Val Met Leu Leu Phe Ala Gly Val Asp Gly
180 185 190

<210> 74

<211> 192

<212> PRT

<213> Homo sapiens

<220>

<223> Individual Isolate: T3

<400> 74

Tyr Glu Val Arg Asn Val Ser Gly Val Tyr Tyr Val Thr Asn Asp Cys
1 5 10 15

Ser Asn Ser Ser Ile Val Tyr Glu Thr Ala Asp Met Ile Met His Thr
20 25 30

Pro Gly Cys Val Pro Cys Val Arg Glu Ser Asn Ser Ser Arg Cys Trp
35 40 45

Val Ala Leu Thr Pro Thr Leu Ala Ala Arg Asn Ala Ser Val Pro Thr
50 55 60

Lys Thr Ile Arg Arg His Val Asp Leu Leu Val Gly Ala Ala Ala Phe
65 70 75 80

Cys Ser Ala Met Tyr Val Gly Asp Leu Cys Gly Ser Val Phe Leu Val
85 90 95

Ser Gln Leu Phe Thr Phe Ser Pro Arg Arg His Glu Thr Val Gln Asp
100 105 110

Cys Asn Cys Ser Ile Tyr Pro Gly His Val Thr Gly His Arg Met Ala
115 120 125

Trp Asp Met Met Met Asn Trp Ser Pro Thr Thr Ala Leu Val Val Ser
130 135 140

Gln Leu Leu Arg Ile Pro Gln Ala Val Val Asp Met Val Ala Gly Ala
145 150 155 160

His Trp Gly Val Leu Ala Gly Leu Ala Tyr Tyr Ser Met Val Gly Asn
165 170 175

Trp Ala Lys Val Leu Ile Val Leu Leu Phe Ala Gly Val Asp Gly
180 185 190

<210> 75
<211> 192
<212> PRT
<213> Homo sapiens

<220>
<223> Individual Isolate: T10

<400> 75
Tyr Glu Val Arg Asn Val Ser Gly Met Tyr His Val Thr Asn Asp Cys
1 5 10 15

Ser Asn Ser Ser Ile Val Phe Glu Ala Ala Asp Leu Ile Met His Thr
20 25 30

Pro Gly Cys Val Pro Cys Val Arg Glu Gly Asn Ser Ser Arg Cys Trp
35 40 45

Val Ala Leu Thr Pro Thr Leu Ala Ala Arg Asn Thr Ser Val Pro Thr
50 55 60

Thr Thr Ile Arg Arg His Val Asp Leu Val Gly Ala Ala Ala Phe
65 70 75 80

Cys Ser Ala Met Tyr Val Gly Asp Leu Cys Gly Ser Val Phe Leu Val
85 90 95

Ser Gln Leu Phe Thr Phe Ser Pro Arg Arg His Glu Thr Leu Gln Asp
100 105 110

Cys Asn Cys Ser Ile Tyr Pro Gly His Leu Ser Gly His Arg Met Ala
115 120 125

Trp Asp Met Met Met Asn Trp Ser Pro Thr Thr Ala Leu Val Val Ser
130 135 140

Gln Leu Leu Arg Ile Pro Gln Ala Val Met Asp Met Val Thr Gly Ala
145 150 155 160

His Trp Gly Val Leu Ala Gly Leu Ala Tyr Tyr Ser Met Ala Gly Asn
165 170 175

Trp Ala Lys Val Leu Ile Val Met Leu Leu Phe Ala Gly Val Asp Gly
180 185 190

<210> 76
<211> 192
<212> PRT
<213> Homo sapiens

<220>
<223> Individual Isolate: US6

<400> 76
Tyr Glu Val Arg Asn Val Ser Gly Met Tyr His Val Thr Asn Asp Cys
1 5 10 15

Ser Asn Ser Ser Ile Val Tyr Glu Ala Ala Asp Met Ile Met His Thr
20 25 30

Pro Gly Cys Val Pro Cys Val Arg Glu Asn Asn Ser Ser Arg Cys Trp
35 40 45

Val Ala Leu Thr Pro Thr Leu Ala Ala Arg Asn Ala Ser Val Pro Thr
50 55 60

Thr Thr Ile Arg Arg His Val Asp Leu Leu Val Gly Ala Ala Thr Phe
65 70 75 80

Cys Ser Ala Met Tyr Val Gly Asp Leu Cys Gly Ser Val Phe Leu Ile
85 90 95

Ser Gln Leu Phe Thr Phe Ser Pro Arg Gln His Glu Thr Val Gln Asp
100 105 110

Cys Asn Cys Ser Ile Tyr Pro Gly His Val Ser Gly His Arg Met Ala
115 120 125

Trp Asp Met Met Asn Trp Ser Pro Thr Ala Ala Leu Val Val Ser
130 135 140

Gln Leu Leu Arg Ile Pro Gln Ala Val Met Asp Met Val Ala Gly Ala

145 150 155 160

His Trp Gly Val Leu Ala Gly Leu Ala Tyr Tyr Ser Met Val Gly Asn
165 170 175

Trp Ala Lys Val Leu Ile Val Leu Leu Phe Ala Gly Val Asp Gly
180 185 190

<210> 77

<211> 192

<212> PRT

<213> Homo sapiens

<220>

<223> Individual Isolate: T2

<400> 77

Ala Gln Val Arg Asn Thr Ser Arg Gly Tyr Met Val Thr Asn Asp Cys
1 5 10 15

Ser Asn Glu Ser Ile Thr Trp Gln Leu Gln Ala Ala Val Leu His Val
20 25 30

Pro Gly Cys Ile Pro Cys Glu Arg Leu Gly Asn Thr Ser Arg Cys Trp
35 40 45

Ile Pro Val Thr Pro Asn Val Ala Val Arg Gln Pro Gly Ala Leu Thr
50 55 60

Gln Gly Leu Arg Thr His Ile Asp Met Val Val Met Ser Ala Thr Leu
65 70 75 80

Cys Ser Ala Leu Tyr Val Gly Asp Leu Cys Gly Gly Val Met Leu Ala
85 90 95

Ala Gln Met Phe Ile Val Ser Pro Arg Arg His Trp Phe Val Gln Glu
100 105 110

Cys Asn Cys Ser Ile Tyr Pro Gly Thr Ile Thr Gly His Arg Met Ala
115 120 125

Trp Asp Met Met Met Asn Trp Ser Pro Thr Ala Thr Met Ile Leu Ala
130 135 140

Tyr Ala Met Arg Val Pro Glu Val Ile Ile Asp Ile Ile Gly Gly Ala
145 150 155 160

His Trp Gly Val Met Phe Gly Leu Ala Tyr Phe Ser Met Gln Gly Ala
165 170 175

Trp Ala Lys Val Ile Val Ile Leu Leu Ala Ala Gly Val Asp Ala
180 185 190

<210> 78
<211> 192
<212> PRT
<213> Homo sapiens

<220>
<223> Individual Isolate: T4

<400> 78
Ala Gln Val Lys Asn Thr Thr Asn Ser Tyr Met Val Thr Asn Asp Cys
1 5 10 15

Ser Asn Asp Ser Ile Thr Trp Gln Leu Gln Ala Ala Val Leu His Val
20 25 30

Pro Gly Cys Val Pro Cys Glu Lys Thr Gly Asn Thr Ser Arg Cys Trp
35 40 45

Ile Pro Val Ser Pro Asn Val Ala Val Arg Gln Pro Gly Ala Leu Thr
50 55 60

Gln Gly Leu Arg Thr His Ile Asp Met Val Val Met Ser Ala Thr Leu
65 70 75 80

Cys Ser Ala Leu Tyr Val Gly Asp Leu Cys Gly Gly Val Met Leu Ala
85 90 95

Ala Gln Met Phe Ile Val Ser Pro Gln His His Trp Phe Val Gln Asp
100 105 110

Cys Asn Cys Ser Ile Tyr Pro Gly Thr Ile Thr Gly His Arg Met Ala
115 120 125

Trp Asp Met Met Asn Trp Ser Pro Thr Ala Thr Met Ile Leu Ala
130 135 140

Tyr Ala Met Arg Val Pro Glu Val Ile Leu Asp Ile Val Ser Gly Ala
145 150 155 160

His Trp Gly Val Met Phe Gly Leu Ala Tyr Phe Ser Met Gln Gly Ala
165 170 175

Trp Ala Lys Val Val Val Ile Leu Leu Ala Ala Gly Val Asp Ala
180 185 190

<210> 79

<211> 192

<212> PRT

<213> Homo sapiens

<220>

<223> Individual Isolate: T9

<400> 79

Ala Glu Val Lys Asn Thr Ser Thr Ser Tyr Met Val Thr Asn Asp Cys
1 5 10 15

Ser Asn Asp Ser Ile Thr Trp Gln Leu Gln Ala Ala Val Leu His Val
20 25 30

Pro Gly Cys Val Pro Cys Glu Arg Val Gly Asn Ala Ser Arg Cys Trp
35 40 45

Ile Pro Val Ser Pro Asn Val Ala Val Gln Arg Pro Gly Ala Leu Thr
50 55 60

Gln Gly Leu Arg Thr His Ile Asp Met Val Val Met Ser Ala Thr Leu
65 70 75 80

Cys Ser Ala Leu Tyr Val Gly Asp Leu Cys Gly Gly Val Met Leu Ala
85 90 95

Ala Gln Met Phe Ile Ile Ser Pro Gln His His Trp Phe Val Gln Glu
100 105 110

Cys Asn Cys Ser Ile Tyr Pro Gly Thr Ile Thr Gly His Arg Met Ala
115 120 125

Trp Asp Met Met Met Asn Trp Ser Pro Thr Thr Thr Met Ile Leu Ala

130	135	140
Tyr Ala Met Arg Val Pro Glu Val Ile Ile Asp Ile Ile Ser Gly Ala		
145	150	155
His Trp Gly Val Met Phe Gly Leu Ala Tyr Phe Ser Met Gln Gly Ala		
165	170	175
Trp Ala Lys Val Val Val Ile Leu Leu Leu Thr Ala Gly Val Asp Ala		
180	185	190
<210> 80		
<211> 192		
<212> PRT		
<213> Homo sapiens		
<220>		
<223> Individual Isolate: US10		
<400> 80		
Val Gln Val Lys Asn Thr Ser Thr Ser Tyr Met Val Thr Asn Asp Cys		
1	5	10
15		
Ser Asn Asp Ser Ile Thr Trp Gln Leu Glu Ala Ala Val Leu His Val		
20	25	30
Pro Gly Cys Val Pro Cys Glu Lys Val Gly Asn Thr Ser Arg Cys Trp		
35	40	45
Ile Pro Val Ser Pro Asn Val Ala Val Gln Arg Pro Gly Ala Leu Thr		
50	55	60
Gln Gly Leu Arg Thr His Ile Asp Met Val Val Met Ser Ala Thr Leu		
65	70	75
80		
Cys Ser Ala Leu Tyr Val Gly Asp Phe Cys Gly Gly Met Met Leu Ala		
85	90	95
Ala Gln Met Phe Ile Val Ser Pro Arg His His Ser Phe Val Gln Glu		
100	105	110
Cys Asn Cys Ser Ile Tyr Pro Gly Thr Ile Thr Gly His Arg Met Ala		
115	120	125

Trp Asp Met Met Asn Trp Ser Pro Thr Ala Thr Leu Ile Leu Ala
130 135 140

Tyr Val Met Arg Val Pro Glu Val Ile Ile Asp Ile Ile Ser Gly Ala
145 150 155 160

His Trp Gly Val Leu Phe Gly Leu Ala Tyr Phe Ser Met Gln Gly Ala
165 170 175

Trp Ala Lys Val Val Val Ile Leu Leu Ala Ala Gly Val Asp Ala
180 185 190

<210> 81

<211> 192

<212> PRT

<213> Homo sapiens

<220>

<223> Individual Isolate: DK8

<400> 81

Val Glu Val Arg Asn Ile Ser Ser Ser Tyr Tyr Ala Thr Asn Asp Cys
1 5 10 15

Ser Asn Asn Ser Ile Thr Trp Gln Leu Thr Asp Ala Val Leu His Leu
20 25 30

Pro Gly Cys Val Pro Cys Glu Asn Asp Asn Gly Thr Leu Arg Cys Trp
35 40 45

Ile Gln Val Thr Pro Asn Val Ala Val Lys His Arg Gly Ala Leu Thr
50 55 60

His Asn Leu Arg Thr His Val Asp Val Ile Val Met Ala Ala Thr Val
65 70 75 80

Cys Ser Ala Leu Tyr Val Gly Asp Val Cys Gly Ala Val Met Ile Val
85 90 95

Ser Gln Ala Leu Ile Ile Ser Pro Glu Arg His Asn Phe Thr Gln Glu
100 105 110

Cys Asn Cys Ser Ile Tyr Gln Gly His Ile Thr Gly His Arg Met Ala
115 120 125

Trp Asp Met Met Leu Asn Trp Ser Pro Thr Leu Thr Met Ile Leu Ala
130 135 140

Tyr Ala Ala Arg Val Pro Glu Leu Ala Leu Gln Val Val Phe Gly Gly
145 150 155 160

His Trp Gly Val Val Phe Gly Leu Ala Tyr Phe Ser Met Gln Gly Ala
165 170 175

Trp Ala Lys Val Ile Ala Ile Leu Leu Val Ala Gly Val Asp Ala
180 185 190

<210> 82

<211> 192

<212> PRT

<213> Homo sapiens

<220>

<223> Individual Isolate: DK11

<400> 82

Val Glu Val Arg Asn Thr Ser Ser Ser Tyr Tyr Ala Thr Asn Asp Cys
1 5 10 15

Ser Asn Asn Ser Ile Thr Trp Gln Leu Thr Asn Ala Val Leu His Leu
20 25 30

Pro Gly Cys Val Pro Cys Glu Asn Asp Asn Gly Thr Leu His Cys Trp
35 40 45

Ile Gln Val Thr Pro Asn Val Ala Val Lys His Arg Gly Ala Leu Thr
50 55 60

His Asn Leu Arg Ala His Ile Asp Met Ile Val Met Ala Ala Thr Val
65 70 75 80

Cys Ser Ala Leu Tyr Val Gly Asp Val Cys Gly Ala Val Met Ile Val
85 90 95

Ser Gln Ala Phe Ile Val Ser Pro Glu His His His Phe Thr Gln Glu
100 105 110

Cys Asn Cys Ser Ile Tyr Gln Gly His Ile Thr Gly His Arg Met Ala

115

120

125

Trp Asp Met Met Leu Asn Trp Ser Pro Thr Leu Thr Met Ile Leu Ala
130 135 140

Tyr Ala Ala Arg Val Pro Glu Leu Val Leu Glu Val Val Phe Gly Gly
145 150 155 160

His Trp Gly Val Val Phe Gly Leu Ala Tyr Phe Ser Met Gln Gly Ala
165 170 175

Trp Ala Lys Val Ile Ala Ile Leu Leu Leu Val Ala Gly Val Asp Ala
180 185 190

<210> 83

<211> 192

<212> PRT

<213> Homo sapiens

<220>

<223> Individual Isolate: SW3

<400> 83

Val Glu Val Arg Asn Ile Ser Ser Ser Tyr Tyr Ala Thr Asn Asp Cys
1 5 10 15

Ser Asn Ser Ser Ile Thr Trp Gln Leu Thr Asn Ala Val Leu His Leu
20 25 30

Pro Gly Cys Val Pro Cys Glu Asn Asp Asn Gly Thr Leu His Cys Trp
35 40 45

Ile Gln Val Thr Pro Asn Val Ala Val Lys His Arg Gly Ala Leu Thr
50 55 60

His Asn Leu Arg Ala His Val Asp Met Ile Val Met Ala Ala Thr Val
65 70 75 80

Cys Ser Ala Leu Tyr Val Gly Asp Met Cys Gly Ala Val Met Ile Val
85 90 95

Ser Gln Ala Phe Ile Ile Ser Pro Glu Arg His Asn Phe Thr Gln Glu
100 105 110

Cys Asn Cys Ser Ile Tyr Gln Gly Arg Ile Thr Gly His Arg Met Ala
115 120 125

Trp Asp Met Met Leu Asn Trp Ser Pro Thr Leu Thr Met Ile Leu Ala
130 135 140

Tyr Ala Ala Arg Val Pro Glu Leu Val Leu Glu Val Val Phe Gly Gly
145 150 155 160

His Trp Gly Val Val Phe Gly Leu Ala Tyr Phe Ser Met Gln Gly Ala
165 170 175

Trp Ala Lys Val Ile Ala Ile Leu Leu Leu Val Ala Gly Val Asp Ala
180 185 190

<210> 84

<211> 192

<212> PRT

<213> Homo sapiens

<220>

<223> Individual Isolate: T8

<400> 84

Val Glu Val Arg Asn Thr Ser Phe Ser Tyr Tyr Ala Thr Asn Asp Cys
1 5 10 15

Ser Asn Asn Ser Ile Thr Trp Gln Leu Thr Asn Ala Val Leu His Leu
20 25 30

Pro Gly Cys Val Pro Cys Glu Asn Asp Asn Gly Thr Leu Arg Cys Trp
35 40 45

Ile Gln Val Thr Pro Asn Val Ala Val Lys His Arg Gly Ala Leu Thr
50 55 60

His Asn Leu Arg Thr His Val Asp Val Ile Val Met Ala Ala Thr Val
65 70 75 80

Cys Ser Ala Leu Tyr Val Gly Asp Val Cys Gly Ala Val Met Ile Ala
85 90 95

Ser Gln Ala Phe Ile Ile Ser Pro Glu Arg His Asn Phe Thr Gln Glu
100 105 110

Cys Asn Cys Ser Ile Tyr Gln Gly His Ile Thr Gly His Arg Met Ala
115 120 125

Trp Asp Met Met Leu Asn Trp Ser Pro Thr Leu Thr Met Ile Leu Ala
130 135 140

Tyr Ala Ala Arg Val Pro Glu Leu Val Leu Glu Val Val Phe Gly Gly
145 150 155 160

His Trp Gly Val Val Phe Gly Leu Ala Tyr Phe Ser Met Gln Gly Ala
165 170 175

Trp Ala Lys Val Ile Ala Ile Leu Leu Val Ala Gly Val Asp Ala
180 185 190

<210> 85

<211> 192

<212> PRT

<213> Homo sapiens

<220>

<223> Individual Isolate: S83

<400> 85

Val Glu Val Lys Asp Thr Gly Asp Ser Tyr Met Pro Thr Asn Asp Cys
1 5 10 15

Ser Asn Ser Ser Ile Val Trp Gln Leu Glu Gly Ala Val Leu His Thr
20 25 30

Pro Gly Cys Val Pro Cys Glu Arg Thr Ala Asn Val Ser Arg Cys Trp
35 40 45

Val Pro Val Ala Pro Asn Leu Ala Ile Ser Gln Pro Gly Ala Leu Thr
50 55 60

Lys Gly Leu Arg Ala His Ile Asp Ile Ile Val Met Ser Ala Thr Val
65 70 75 80

Cys Ser Ala Leu Tyr Val Gly Asp Val Cys Gly Ala Leu Met Leu Ala
85 90 95

Ala Gln Val Val Val Ser Pro Gln His His Thr Phe Val Gln Glu

100 105 110
Cys Asn Cys Ser Ile Tyr Pro Gly Arg Ile Thr Gly His Arg Met Ala
115 120 125

Trp Asp Met Met Met Asn Trp Ser Pro Thr Thr Thr Met Leu Leu Ala
130 135 140

Tyr Leu Val Arg Ile Pro Glu Val Ile Leu Asp Ile Val Thr Gly Gly
145 150 155 160

His Trp Gly Val Met Phe Gly Leu Ala Tyr Phe Ser Met Gln Gly Ser
165 170 175

Trp Ala Lys Val Ile Val Ile Leu Leu Leu Thr Ala Gly Val Glu Ala
180 185 190

<210> 86

<211> 192

<212> PRT

<213> Homo sapiens

<220>

<223> Individual Isolate: DK12

<400> 86

Leu Glu Trp Arg Asn Val Ser Gly Leu Tyr Val Leu Thr Asn Asp Cys
1 5 10 15

Ser Asn Ser Ser Ile Val Tyr Glu Ala Asp Asp Val Ile Leu His Thr
20 25 30

Pro Gly Cys Val Pro Cys Val Gln Asp Gly Asn Thr Ser Thr Cys Trp
35 40 45

Thr Ser Val Thr Pro Thr Val Ala Val Arg Tyr Val Gly Ala Thr Thr
50 55 60

Ala Ser Ile Arg Ser His Val Asp Leu Leu Val Gly Ala Ala Thr Met
65 70 75 80

Cys Ser Ala Leu Tyr Val Gly Asp Val Cys Gly Ala Val Phe Leu Val
85 90 95

Gly Gln Ala Phe Thr Phe Arg Pro Arg Arg His Gln Thr Val Gln Thr
100 105 110

Cys Asn Cys Ser Leu Tyr Pro Gly His Leu Ser Gly His Arg Met Ala
115 120 125

Trp Asp Met Met Met Asn Trp Ser Pro Ala Val Gly Met Val Val Ala
130 135 140

His Val Leu Arg Leu Pro Gln Thr Leu Phe Asp Ile Ile Ala Gly Ala
145 150 155 160

His Trp Gly Ile Met Ala Gly Leu Ala Tyr Tyr Ser Met Gln Gly Asn
165 170 175

Trp Ala Lys Val Ala Ile Ile Met Val Met Phe Ser Gly Val Asp Ala
180 185 190

<210> 87
<211> 192
<212> PRT
<213> Homo sapiens

<220>
<223> Individual Isolate: HK10

<400> 87
Leu Glu Trp Arg Asn Val Ser Gly Leu Tyr Val Leu Thr Asn Asp Cys
1 5 10 15

Pro Asn Ser Ser Ile Val Tyr Glu Ala Asp Asp Val Ile Leu His Thr
20 25 30

Pro Gly Cys Val Pro Cys Val Gln Asp Gly Asn Thr Ser Thr Cys Trp
35 40 45

Thr Ser Val Thr Pro Thr Val Ala Val Arg Tyr Val Gly Ala Thr Thr
50 55 60

Ala Ser Ile Arg Ser His Val Asp Leu Leu Val Gly Ala Ala Thr Met
65 70 75 80

Cys Ser Ala Leu Tyr Val Gly Asp Met Cys Gly Ala Val Phe Leu Val
85 90 95

Gly Gln Ala Phe Thr Phe Arg Pro Arg Arg His Gln Thr Val Gln Thr
100 105 110

Cys Asn Cys Ser Leu Tyr Pro Gly His Leu Ser Gly His Arg Met Ala
115 120 125

Trp Asp Met Met Met Asn Trp Ser Pro Ala Val Gly Met Val Val Ala
130 135 140

His Val Leu Arg Leu Pro Gln Thr Leu Phe Asp Ile Ile Ala Gly Ala
145 150 155 160

His Trp Gly Ile Leu Ala Gly Leu Ala Tyr Tyr Ser Met Gln Gly Asn
165 170 175

Trp Ala Lys Val Ala Ile Ile Met Val Met Phe Ser Gly Val Asp Ala
180 185 190

<210> 88

<211> 192

<212> PRT

<213> Homo sapiens

<220>

<223> Individual Isolate: S2

<400> 88

Leu Glu Trp Arg Asn Thr Ser Gly Leu Tyr Val Leu Thr Asn Asp Cys
1 5 10 15

Ser Asn Ser Ser Ile Val Tyr Glu Ala Asp Asp Val Ile Leu His Thr
20 25 30

Pro Gly Cys Val Pro Cys Val Gln Asp Gly Asn Thr Ser Thr Cys Trp
35 40 45

Thr Pro Val Thr Pro Thr Val Ala Val Arg Tyr Val Gly Ala Thr Thr
50 55 60

Ala Ser Ile Arg Ser His Val Asp Leu Leu Val Gly Ala Ala Thr Met
65 70 75 80

Cys Ser Ala Leu Tyr Val Gly Asp Met Cys Gly Ala Val Phe Leu Val

85

90

95

Gly Gln Ala Phe Thr Phe Arg Pro Arg Arg His Gln Thr Val Gln Thr
100 105 110

Cys Asn Cys Ser Leu Tyr Pro Gly His Leu Ser Gly His Arg Met Ala
115 120 125

Trp Asp Met Met Asn Trp Ser Pro Ala Val Gly Met Val Val Ala
130 135 140

His Val Leu Arg Leu Pro Gln Thr Val Phe Asp Ile Ile Ala Gly Ala
145 150 155 160

His Trp Gly Ile Leu Ala Gly Leu Ala Tyr Tyr Ser Met Gln Gly Asn
165 170 175

Trp Ala Lys Val Ala Ile Ile Met Val Met Phe Ser Gly Val Asp Ala
180 185 190

<210> 89

<211> 192

<212> PRT

<213> Homo sapiens

<220>

<223> Individual Isolate: S52

<400> 89

Leu Glu Trp Arg Asn Thr Ser Gly Leu Tyr Val Leu Thr Asn Asp Cys
1 5 10 15

Ser Asn Ser Ser Ile Val Tyr Glu Ala Asp Asp Val Ile Leu His Thr
20 25 30

Pro Gly Cys Val Pro Cys Val Gln Asp Gly Asn Thr Ser Met Cys Trp
35 40 45

Thr Pro Val Thr Pro Thr Val Ala Val Arg Tyr Val Gly Ala Thr Thr
50 55 60

Ala Ser Ile Arg Ser His Val Asp Leu Leu Val Gly Ala Ala Thr Leu
65 70 75 80

Cys	Ser	Ala	Leu	Tyr	Val	Gly	Asp	Met	Cys	Gly	Ala	Val	Phe	Leu	Val
85									90						95
Gly	Gln	Ala	Phe	Thr	Phe	Arg	Pro	Arg	Arg	His	Gln	Thr	Val	Gln	Thr
100						105									110
Cys	Asn	Cys	Ser	Leu	Tyr	Pro	Gly	His	Val	Ser	Gly	His	Arg	Met	Ala
115						120									125
Trp	Asp	Met	Met	Met	Asn	Trp	Ser	Pro	Ala	Val	Gly	Met	Val	Val	Ala
130						135						140			
His	Ile	Leu	Arg	Leu	Pro	Gln	Thr	Leu	Phe	Asp	Ile	Leu	Ala	Gly	Ala
145						150					155				160
His	Trp	Gly	Ile	Leu	Ala	Gly	Leu	Ala	Tyr	Tyr	Ser	Met	Gln	Gly	Asn
165						170						175			
Trp	Ala	Lys	Val	Ala	Ile	Val	Met	Ile	Met	Phe	Ser	Gly	Val	Asp	Ala
180						185						190			

<210> 90
 <211> 192
 <212> PRT
 <213> Homo sapiens

<220>
 <223> Individual Isolate: S54

<400> 90
 Leu Glu Trp Arg Asn Thr Ser Gly Leu Tyr Ile Leu Thr Asn Asp Cys
 1 5 10 15

Ser Asn Ser Ser Ile Val Tyr Glu Ala Asp Asp Val Ile Leu His Thr
 20 25 30

Pro Gly Cys Val Pro Cys Val Gln Asp Gly Asn Thr Ser Thr Cys Trp
 35 40 45

Thr Pro Val Thr Pro Thr Val Ala Val Arg Tyr Val Gly Ala Thr Thr
 50 55 60

Ala Ser Ile Arg Ser His Val Asp Leu Leu Val Gly Ala Ala Thr Leu
 65 70 75 80

Cys Ser Ala Leu Tyr Val Gly Asp Met Cys Gly Ala Val Phe Leu Val
85 90 95

Gly Gln Ala Phe Thr Phe Arg Pro Arg Arg His Gln Thr Val Gln Thr
100 105 110

Cys Asn Cys Ser Leu Tyr Pro Gly His Leu Ser Gly His Arg Met Ala
115 120 125

Trp Asp Met Met Met Asn Trp Ser Pro Ala Val Gly Met Val Val Ala
130 135 140

His Ile Leu Arg Leu Pro Gln Thr Leu Phe Asp Ile Leu Ala Gly Ala
145 150 155 160

His Trp Gly Ile Leu Ala Gly Leu Ala Tyr Tyr Ser Met Gln Gly Asn
165 170 175

Trp Ala Lys Val Ala Ile Ile Met Ile Met Phe Ser Gly Val Asp Ala
180 185 190

<210> 91

<211> 192

<212> PRT

<213> Homo sapiens

<220>

<223> Individual Isolate: Z4

<400> 91

Glu His Tyr Arg Asn Ala Ser Gly Ile Tyr His Ile Thr Asn Asp Cys
1 5 10 15

Pro Asn Ser Ser Ile Val Tyr Glu Ala Asp His His Ile Leu His Leu
20 25 30

Pro Gly Cys Val Pro Cys Val Met Thr Gly Asn Thr Ser Arg Cys Trp
35 40 45

Thr Pro Val Thr Pro Thr Val Ala Val Ala His Pro Gly Ala Pro Leu
50 55 60

Glu Ser Phe Arg Arg His Val Asp Leu Met Val Gly Ala Ala Thr Leu

65 70 75 80

Cys Ser Ala Leu Tyr Val Gly Asp Leu Cys Gly Gly Ala Phe Leu Met
85 90 95

Gly Gln Met Ile Thr Phe Arg Pro Arg Arg His Trp Thr Thr Gln Glu
100 105 110

Cys Asn Cys Ser Ile Tyr Thr Gly His Ile Thr Gly His Arg Met Ala
115 120 125

Trp Asp Met Met Asn Trp Ser Pro Thr Thr Thr Leu Leu Leu Ala
130 135 140

Gln Ile Met Arg Val Pro Thr Ala Phe Leu Asp Met Val Ala Gly Gly
145 150 155 160

His Trp Gly Val Leu Ala Gly Leu Ala Tyr Phe Ser Met Gln Gly Asn
165 170 175

Trp Ala Lys Val Val Leu Val Leu Phe Leu Phe Ala Gly Val Asp Ala
180 185 190

<210> 92

<211> 192

<212> PRT

<213> Homo sapiens

<220>

<223> Individual Isolate: Z1

<400> 92

Val His Tyr Arg Asn Ala Ser Gly Val Tyr His Val Thr Asn Asp Cys
1 5 10 15

Pro Asn Thr Ser Ile Val Tyr Glu Thr Glu His His Ile Met His Leu
20 25 30

Pro Gly Cys Val Pro Cys Val Arg Thr Glu Asn Thr Ser Arg Cys Trp
35 40 45

Val Pro Leu Thr Pro Thr Val Ala Ala Pro Tyr Pro Asn Ala Pro Leu
50 55 60

Glu Ser Met Arg Arg His Val Asp Leu Met Val Gly Ala Ala Thr Met
65 70 75 80

Cys Ser Ala Phe Tyr Ile Gly Asp Leu Cys Gly Gly Val Phe Leu Val
85 90 95

Gly Gln Leu Phe Asp Phe Arg Pro Arg Arg His Trp Thr Thr Gln Asp
100 105 110

Cys Asn Cys Ser Ile Tyr Pro Gly His Val Ser Gly His Arg Met Ala
115 120 125

Trp Asp Met Met Met Asn Trp Ser Pro Thr Ser Ala Leu Ile Met Ala
130 135 140

Gln Ile Leu Arg Ile Pro Ser Ile Leu Gly Asp Leu Leu Thr Gly Gly
145 150 155 160

His Trp Gly Val Leu Ala Gly Leu Ala Phe Phe Ser Met Gln Ser Asn
165 170 175

Trp Ala Lys Val Ile Leu Val Leu Phe Leu Phe Ala Gly Val Glu Gly
180 185 190

<210> 93
<211> 192
<212> PRT
<213> Homo sapiens

<220>
<223> Individual Isolate: Z6

<400> 93
Val Asn Tyr Arg Asn Ala Ser Gly Val Tyr His Val Thr Asn Asp Cys
1 5 10 15

Pro Asn Ser Ser Ile Val Tyr Glu Ala Glu His Gln Ile Leu His Leu
20 25 30

Pro Gly Cys Leu Pro Cys Val Arg Val Gly Asn Gln Ser Arg Cys Trp
35 40 45

Val Ala Leu Thr Pro Thr Val Ala Val Ser Tyr Ile Gly Ala Pro Leu
50 55 60

Asp Ser Leu Arg Arg His Val Asp Leu Met Val Gly Ala Ala Thr Val			
65	70	75	80
Cys Ser Ala Leu Tyr Val Gly Asp Leu Cys Gly Gly Ala Phe Leu Val			
85	90	95	
Gly Gln Met Phe Ser Phe Gln Pro Arg Arg His Trp Thr Thr Gln Asp			
100	105	110	
Cys Asn Cys Ser Ile Tyr Ala Gly His Ile Thr Gly His Arg Met Ala			
115	120	125	
Trp Asp Met Met Asn Trp Ser Pro Thr Thr Thr Leu Leu Leu Ala			
130	135	140	
Gln Val Met Arg Ile Pro Ser Thr Leu Val Asp Leu Leu Ala Gly Gly			
145	150	155	160
His Trp Gly Val Leu Val Gly Leu Ala Tyr Phe Ser Met Gln Ala Asn			
165	170	175	
Trp Ala Lys Val Ile Leu Val Leu Phe Leu Phe Ala Gly Val Asp Ala			
180	185	190	

<210> 94
 <211> 192
 <212> PRT
 <213> Homo sapiens

<220>
 <223> Individual Isolate: Z7

<400> 94
 Val Asn Tyr His Asn Ala Ser Gly Val Tyr His Ile Thr Asn Asp Cys
 1 5 10 15

Pro Asn Ser Ser Ile Met Tyr Glu Ala Glu His His Ile Leu His Leu
 20 25 30

Pro Gly Cys Val Pro Cys Val Arg Glu Gly Asn Gln Ser Arg Cys Trp
 35 40 45

Val Ala Leu Thr Pro Thr Val Ala Ala Pro Tyr Ile Gly Ala Pro Leu

50	55	60
Glu Ser Ile Arg Arg His Val Asp Leu Met Val Gly Ala Ala Thr Val		
65	70	75
Cys Ser Ala Leu Tyr Ile Gly Asp Leu Cys Gly Gly Val Phe Leu Val		
85	90	95
Gly Gln Met Phe Ser Phe Gln Pro Arg Arg His Trp Thr Thr Gln Asp		
100	105	110
Cys Asn Cys Ser Ile Tyr Ala Gly His Val Thr Gly His Arg Met Ala		
115	120	125
Trp Asp Met Met Met Asn Trp Ser Pro Thr Thr Thr Leu Val Leu Ala		
130	135	140
Gln Val Met Arg Ile Pro Ser Thr Leu Val Asp Leu Leu Thr Gly Gly		
145	150	155
His Trp Gly Ile Leu Ile Gly Val Ala Tyr Phe Cys Met Gln Ala Asn		
165	170	175
Trp Ala Lys Val Ile Leu Val Leu Phe Leu Tyr Ala Gly Val Asp Ala		
180	185	190

<210> 95
 <211> 192
 <212> PRT
 <213> Homo sapiens

<220>
 <223> Individual Isolate: DK13

<400> 95
 Tyr Asn Tyr Arg Asn Ser Ser Gly Val Tyr His Val Thr Asn Asp Cys
 1 5 10 15

Pro Asn Ser Ser Ile Val Tyr Glu Thr Asp Tyr His Ile Leu His Leu
 20 25 30

Pro Gly Cys Val Pro Cys Val Arg Glu Gly Asn Lys Ser Thr Cys Trp
 35 40 45

Val	Ser	Leu	Thr	Pro	Thr	Val	Ala	Ala	Gln	His	Leu	Asn	Ala	Pro	Leu
50						55				60					
Glu	Ser	Leu	Arg	Arg	His	Val	Asp	Leu	Met	Val	Gly	Gly	Ala	Thr	Leu
65						70			75				80		
Cys	Ser	Ala	Leu	Tyr	Ile	Gly	Asp	Val	Cys	Gly	Gly	Val	Phe	Leu	Val
					85				90				95		
Gly	Gln	Leu	Phe	Thr	Phe	Gln	Pro	Arg	Arg	His	Trp	Thr	Thr	Gln	Asp
					100			105				110			
Cys	Asn	Cys	Ser	Ile	Tyr	Thr	Gly	His	Ile	Thr	Gly	His	Arg	Met	Ala
					115			120				125			
Trp	Asp	Met	Met	Met	Asn	Trp	Ser	Pro	Thr	Ala	Thr	Leu	Val	Leu	Ala
					130			135			140				
Gln	Leu	Met	Arg	Ile	Pro	Gly	Ala	Met	Val	Asp	Leu	Leu	Ala	Gly	Gly
					145			150			155			160	
His	Trp	Gly	Ile	Leu	Val	Gly	Ile	Ala	Tyr	Phe	Ser	Met	Gln	Ala	Asn
					165			170				175			
Trp	Ala	Lys	Val	Ile	Leu	Val	Leu	Phe	Leu	Phe	Ala	Gly	Val	Asp	Ala
					180			185				190			

<210> 96
 <211> 192
 <212> PRT
 <213> Homo sapiens

<220>
 <223> Individual Isolate: SA1

<400> 96
 Val Pro Tyr Arg Asn Ala Ser Gly Val Tyr His Val Thr Asn Asp Cys
 1 5 10 15

Pro Asn Ser Ser Ile Val Tyr Glu Ala Asp Ser Leu Ile Leu His Ala
 20 25 30

Pro Gly Cys Val Pro Cys Val Arg Gln Asp Asn Val Ser Arg Cys Trp
 35 40 45

Val Gln Ile Thr Pro Thr Leu Ser Ala Pro Thr Phe Gly Ala Val Thr			
50	55	60	
Ala Pro Leu Arg Arg Ala Val Asp Tyr Leu Ala Gly Gly Ala Ala Leu			
65	70	75	80
Cys Ser Ala Leu Tyr Val Gly Asp Ala Cys Gly Ala Val Phe Leu Val			
85	90	95	
Gly Gln Met Phe Thr Tyr Arg Pro Arg Gln His Thr Thr Val Gln Asp			
100	105	110	
Cys Asn Cys Ser Ile Tyr Ser Gly His Ile Thr Gly His Arg Met Ala			
115	120	125	
Trp Asp Met Met Asn Trp Ser Pro Thr Thr Ala Leu Leu Met Ala			
130	135	140	
Gln Met Leu Arg Ile Pro Gln Val Val Ile Asp Ile Ile Ala Gly Gly			
145	150	155	160
His Trp Gly Val Leu Phe Ala Ala Ala Tyr Phe Ala Ser Ala Ala Asn			
165	170	175	
Trp Ala Lys Val Val Leu Val Leu Phe Leu Phe Ala Gly Val Asp Gly			
180	185	190	

<210> 97
 <211> 192
 <212> PRT
 <213> Homo sapiens

<220>
 <223> Individual Isolate: SA4

<400> 97
 Val Pro Tyr Arg Asn Ala Ser Gly Val Tyr His Val Thr Asn Asp Cys
 1 5 10 15

Pro Asn Ser Ser Ile Val Tyr Glu Ala Asp Asn Leu Ile Leu His Ala
 20 25 30

Pro Gly Cys Val Pro Cys Val Arg Gln Asp Asn Val Ser Lys Cys Trp

35	40	45
Val Gln Ile Thr Pro Thr Leu Ser Ala Pro Asn Leu Gly Ala Val Thr		
50	55	60
Ala Pro Leu Arg Arg Ala Val Asp Tyr Leu Ala Gly Gly Ala Ala Leu		
65	70	75
Cys Ser Ala Leu Tyr Val Gly Asp Ala Cys Gly Ala Val Phe Leu Val		
85	90	95
Gly Gln Met Phe Thr Tyr Arg Pro Arg Gln His Thr Thr Val Gln Asp		
100	105	110
Cys Asn Cys Ser Ile Tyr Ser Gly His Ile Thr Gly His Arg Met Ala		
115	120	125
Trp Asp Met Met Met Asn Trp Ser Pro Thr Thr Ala Leu Leu Met Ala		
130	135	140
Gln Leu Leu Arg Ile Pro Gln Val Val Ile Asp Ile Ile Ala Gly Gly		
145	150	155
160		
His Trp Gly Val Leu Phe Ala Ala Ala Tyr Phe Ala Ser Ala Ala Asn		
165	170	175
Trp Ala Lys Val Ile Leu Val Leu Phe Leu Phe Ala Gly Val Asp Ala		
180	185	190

<210> 98
 <211> 192
 <212> PRT
 <213> Homo sapiens

<220>
 <223> Individual Isolate: SA5

<400> 98
 Val Pro Tyr Arg Asn Ala Ser Gly Val Tyr His Val Thr Asn Asp Cys
 1 5 10 15
 Pro Asn Ser Ser Ile Val Tyr Glu Ala Asp Asn Leu Ile Leu His Ala
 20 25 30

Pro	Gly	Cys	Val	Pro	Cys	Val	Lys	Glu	Gly	Asn	Val	Ser	Arg	Cys	Trp
35							40						45		
Val	Gln	Ile	Thr	Pro	Thr	Leu	Ser	Ala	Pro	Asn	Leu	Gly	Ala	Val	Thr
50						55					60				
Ala	Pro	Leu	Arg	Arg	Val	Val	Asp	Tyr	Leu	Ala	Gly	Gly	Ala	Ala	Leu
65					70				75				80		
Cys	Ser	Ala	Leu	Tyr	Val	Gly	Asp	Ala	Cys	Gly	Ala	Val	Phe	Leu	Val
85						90						95			
Gly	Gln	Met	Phe	Thr	Tyr	Arg	Pro	Arg	Gln	His	Thr	Thr	Val	Gln	Asp
100						105					110				
Cys	Asn	Cys	Ser	Ile	Tyr	Ser	Gly	His	Ile	Thr	Gly	His	Arg	Met	Ala
115						120				125					
Trp	Asp	Met	Met	Met	Asn	Trp	Ser	Pro	Thr	Thr	Ala	Leu	Val	Met	Ala
130						135				140					
Gln	Val	Leu	Arg	Ile	Pro	Gln	Val	Val	Ile	Asp	Ile	Ile	Ala	Gly	Gly
145						150				155			160		
His	Trp	Gly	Val	Leu	Phe	Ala	Val	Ala	Tyr	Phe	Ala	Ser	Ala	Ala	Asn
165						170					175				
Trp	Ala	Lys	Val	Val	Leu	Val	Phe	Leu	Phe	Ala	Gly	Val	Asp	Gly	
180						185					190				

<210> 99
 <211> 192
 <212> PRT
 <213> Homo sapiens

<220>
 <223> Individual Isolate: SA6

<400> 99
 Val Pro Tyr Arg Asn Ala Ser Gly Val Tyr His Val Thr Asn Asp Cys
 1 5 10 15

Pro Asn Ser Ser Ile Val Tyr Glu Ala Asp Asp Leu Ile Leu His Ala
 20 25 30

Pro	Gly	Cys	Val	Pro	Cys	Val	Arg	Lys	Asp	Asn	Val	Ser	Arg	Cys	Trp
35															45
Val	His	Ile	Thr	Pro	Thr	Leu	Ser	Ala	Pro	Ser	Leu	Gly	Ala	Val	Thr
50															60
Ala	Pro	Leu	Arg	Arg	Ala	Val	Asp	Tyr	Leu	Ala	Gly	Gly	Ala	Ala	Leu
65															80
Cys	Ser	Ala	Leu	Tyr	Val	Gly	Asp	Val	Cys	Gly	Ala	Leu	Phe	Leu	Val
85															95
Gly	Gln	Met	Phe	Thr	Tyr	Arg	Pro	Arg	Gln	His	Ala	Thr	Val	Gln	Asp
100															110
Cys	Asn	Cys	Ser	Ile	Tyr	Ser	Gly	His	Ile	Thr	Gly	His	Arg	Met	Ala
115															125
Trp	Asp	Met	Met	Met	Asn	Trp	Ser	Pro	Ala	Thr	Ala	Leu	Val	Met	Ala
130															140
Gln	Met	Leu	Arg	Ile	Pro	Gln	Val	Val	Ile	Asp	Ile	Ile	Ala	Gly	Gly
145															160
His	Trp	Gly	Val	Leu	Phe	Ala	Ala	Ala	Tyr	Phe	Ala	Ser	Ala	Ala	Asn
165															175
Trp	Ala	Lys	Val	Val	Leu	Val	Leu	Phe	Leu	Phe	Ala	Gly	Val	Asp	Ala
180															190

<210> 100
 <211> 192
 <212> PRT
 <213> Homo sapiens

<220>
 <223> Individual Isolate: SA7

<400> 100
 Val Pro Tyr Arg Asn Ala Ser Gly Val Tyr His Val Thr Asn Asp Cys
 1 5 10 15

Pro Asn Ser Ser Ile Val Tyr Glu Ala Asp Asn Leu Ile Leu His Ala

20	25	30
Pro Gly Cys Val Pro Cys Val Arg Gln Asn Asn Val Ser Arg Cys Trp		
35	40	45
Val Gln Ile Thr Pro Thr Leu Ser Ala Pro Asn Leu Gly Ala Val Thr		
50	55	60
Ala Pro Leu Arg Arg Ala Val Asp Tyr Leu Ala Gly Gly Ala Ala Leu		
65	70	75
Cys Ser Ala Leu Tyr Val Gly Asp Ala Cys Gly Ala Val Phe Leu Val		
85	90	95
Gly Gln Met Phe Ser Tyr Arg Pro Arg Gln His Thr Thr Val Gln Asp		
100	105	110
Cys Asn Cys Ser Ile Tyr Ser Gly His Ile Thr Gly His Arg Met Ala		
115	120	125
Trp Asp Met Met Met Asn Trp Ser Pro Thr Thr Ala Leu Val Met Ala		
130	135	140
Gln Leu Leu Arg Ile Pro Gln Val Val Ile Asp Ile Ile Ala Gly Gly		
145	150	155
His Trp Gly Val Leu Phe Ala Ala Ala Tyr Phe Ala Ser Ala Ala Asn		
165	170	175
Trp Ala Lys Val Val Leu Val Leu Phe Leu Phe Ala Gly Val Asp Ala		
180	185	190

<210> 101
<211> 192
<212> PRT
<213> *Homo sapiens*

<220>
<223> Individual Isolate: SA13

<400> 101
Val Pro Tyr Arg Asn Ala Ser Gly Val Tyr His Val Thr Asn Asp Cys
1 5 10 15

Pro	Asn	Ser	Ser	Ile	Val	Tyr	Glu	Ala	Asp	Asp	Leu	Ile	Leu	His	Ala
20														30	
Pro	Gly	Cys	Val	Pro	Cys	Val	Arg	Gln	Gly	Asn	Val	Ser	Arg	Cys	Trp
35														45	
Val	Gln	Ile	Thr	Pro	Thr	Leu	Ser	Ala	Pro	Ser	Leu	Gly	Ala	Val	Thr
50														60	
Ala	Pro	Leu	Arg	Arg	Ala	Val	Asp	Tyr	Leu	Ala	Gly	Gly	Ala	Ala	Leu
65														80	
Cys	Ser	Ala	Leu	Tyr	Val	Gly	Asp	Ala	Cys	Gly	Ala	Val	Phe	Leu	Val
85														95	
Gly	Gln	Met	Phe	Thr	Tyr	Ser	Pro	Arg	Arg	His	Asn	Val	Val	Gln	Asp
100														110	
Cys	Asn	Cys	Ser	Ile	Tyr	Ser	Gly	His	Ile	Thr	Gly	His	Arg	Met	Ala
115														125	
Trp	Asp	Met	Met	Met	Asn	Trp	Ser	Pro	Thr	Thr	Ala	Leu	Val	Met	Ala
130														140	
Gln	Leu	Leu	Arg	Ile	Pro	Gln	Val	Val	Ile	Asp	Ile	Ile	Ala	Gly	Ala
145														160	
His	Trp	Gly	Val	Leu	Phe	Ala	Ala	Ala	Tyr	Tyr	Ala	Ser	Ala	Ala	Asn
165														175	
Trp	Ala	Lys	Val	Val	Leu	Val	Leu	Phe	Leu	Phe	Ala	Gly	Val	Asp	Ala
180														190	

<210> 102

<211> 192

<212> PRT

<213> Homo sapiens

<220>

<223> Individual Isolate: HK2

<400> 102

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Pro Asn Ser Ser Ile Val Leu Glu Ala Asp Ala Met Ile Leu His Leu
20 25 30

Pro Gln Cys Leu Pro Cys Val Arg Val Asp Asp Arg Ser Thr Cys Trp
35 40 45

His Ala Val Thr Pro Thr Leu Ala Ile Pro Asn Ala Ser Thr Pro Ala
50 55 60

Thr Gln Phe Arg Arg His Val Asp Leu Leu Ala Gln Ala Ala Val Val
65 70 75 80

Cys Ser Ser Leu Tyr Ile Gln Asp Leu Cys Gln Ser Leu Phe Leu Ala
85 90 95

Gln Gln Leu Phe Thr Phe Gln Pro Arg Arg His Trp Thr Val Gln Asp
100 105 110

Cys Asn Cys Ser Ile Tyr Thr Gln His Val Thr Gln His Arg Met Ala
115 120 125

Trp Asp Met Met Met Asn Trp Ser Pro Thr Thr Thr Leu Val Leu Ser
130 135 140

Ser Ile Leu Arg Val Pro Glu Ile Cys Ala Ser Val Ile Phe Gln Gln
145 150 155 160

His Trp Gln Ile Leu Leu Ala Val Ala Tyr Phe Gln Met Ala Gln Asn
165 170 175

Trp Leu Lys Val Leu Ala Val Leu Phe Leu Phe Ala Gln Val Glu Ala
180 185 190

<210> 103

<211> 573

<212> DNA

<213> Homo sapiens

<220>

<223> Individual Isolate: DK7

<400> 103

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tacccttggc ccctctatgg caatgagggc tgccgggtggg cgggatggct cctgtctccc 300
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ggcgccccctc ttggaggcgca tgccagggcc ctggcgcatg gcgtccgggt tctggaagac 480
ggcgtgaact atgcaacagg gaaccttcctt ggttgcctt tctctatctt ccttggcc 540
ctgctcttgc ctgactgt gcccgttca gcc 573

<210> 104

<211> 573

<212> DNA

<213> Homo sapiens

<220>

<223> Individual Isolate: US11

<400> 104

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ggccctagat tgggtgtcg cgccgcgagg aagacttccg agcggtcgca acctcgaggt 180
agacgtcagc ctatccccaa ggcacgtcg cccgagggca ggacctggc tcagcccggg 240
tacccttggc ccctctatgg caatgagggc tgccgggtggg cgggatggct cctgtctccc 300
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ggcgccccctc tcggaggcgca tgccagggcc ctggcgcatg gcgtccgggt tctggaagac 480
ggcgtgaact atgcaacagg gaaccttcctt ggttgcctt tctctatctt ccttggcc 540
ctgctcttgc ctgactgt gcccgttca gcc 573

<210> 105

<211> 573

<212> DNA

<213> Homo sapiens

<220>

<223> Individual Isolate: S14

<400> 105

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gacgtcaagt tcccgggtgg cggtcagatc gttgggtggag tttacttgtt gccgcgcagg 120
ggccctagat tgggtgtcg cgccgcgagg aagacttccg agcggtcgca acctcgaggt 180
agacgtcagc ctatccccaa ggcacgtcg cccgagggca ggacctggc tcagcccggg 240
tacccttggc ccctctatgg caatgagggc tgccgggtggg cgggatggct cctgtctccc 300
cgtggctctc ggcctagctg gggccccaca gaccccccggc gtaggtcgca caatttgggt 360
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ggcgcccccc tcgggggcgc tgccagggcc ctggcgcatg gcgtccgggt tctggaagac 480
ggcgtgaact atgcaacagg gaaccttcct gggtgctctt tctctatctt cctcctagcc 540
ctgctttctt gcctgactgt gcccgttca gcc 573

<210> 106
<211> 573
<212> DNA
<213> Homo sapiens

<220>
<223> Individual Isolate: SW1

<400> 106
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gacgtcaagt tcccgggtgg cggtcagatc gttgggtggag tttacttgg tccgcgcagg 120
ggccctagat tgggtgtgcg cgcgacgagg aagacttccg agcggtcgca acctcgaggt 180
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cgtggctctc ggcttagctg gggccctaca gaccccccggc gtaggtcgca caatttgggt 360
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ggcgccccctc ttggaggcgc tgccagggcc ctggcgcatg gcgtccgggt tctggaagac 480
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ctgctttctt gcctgacagt gcccgttca gcc 573

<210> 107
<211> 573
<212> DNA
<213> Homo sapiens

<220>
<223> Individual Isolate: S18

<400> 107
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gacgttaagt tcccgggtgg cggtcagatc gttgggtggag tttacttgg tccgcgcagg 120
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tacccttggc ccctctatgg caatgagggc tgccggatggg cggtatggct cctgtcccc 300
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ggcgccccctc tcggaggcgc tgccagggcc ctggcgcatg gcgtccgggt tctggaagac 480
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ctgctctctt gtctgactgt gcccgttca gct 573

<210> 108

<211> 573
<212> DNA
<213> Homo sapiens

<220>
<223> Individual Isolate: DR4

<400> 108
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gacgtcaagt tcccgggtgg cggtcagatc gttggtggag tttacttgg 573
ggccctagat tgggtgtgcg cgcgacgagg aagacttccg agcggtcgca acctcgaggt 180
agacgtcagc ctatccccaa ggccgtcg 573
cccgaggca ggacctgggc tcagccccc 240
tacccttggc ccctctatgg caatgagggc tgccgggtggg cggatggct cctgtcccc 300
cgtggctctc ggcctagctg gggccccaca gaccccccgc gtaggtcg 573
aaggtcatcg acaccctcac gtgcggcttc gccgaccta tgggtacat cccgctcg 420
ggcccccccc ttgggggcgc tgccagggcc ctggcgcatg gcgtccgag 573
ggcgtgaact atgcaacagg gaatcttcc 573
ttgctcttgc gcttggaccgt gcccgcatacg gcc 573

<210> 109
<211> 573
<212> DNA
<213> Homo sapiens

<220>
<223> Individual Isolate: SA10

<400> 109
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gacgtcaagt tcccgggcgg tgggtcagatc gttggtggag tctatctgg 573
ggccccaggt tgggtgtgcg cgcgacgagg aagacttccg agcggtcgca acctcg 573
aggcgacaac ctatccccaa ggctcgccag cccgaggca ggacctgggc ccagccccc 240
tacccttggc ccctctatgg caatgagggc ttgggtggg caggatggct cctgtcaccc 300
cgtggctctc ggcctagttg gggccccacg gaccccccgc gtaggtcg 573
aaggtcatcg ataccctcac atgcggcttc gccgaccta tgggtacat tccgctcg 420
ggcgccccctt tagggggcgc tgccagggcc ttggcgcatg gcgtccgggt tctgg 573
ggcgtgaact atgcaacagg gaatttgc 573
ttgctgtccat gtttaaccat cccagcttcc gct 573

<210> 110
<211> 573
<212> DNA
<213> Homo sapiens

<220>
<223> Individual Isolate: S45

<400> 110
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gacgtcaagt tcccgggtgg cggtcagatc gttgggtggag tttacctgtt gccgcgcagg 120
ggccccaggt tgggtgtgcg cgcgactagg aagacttccg agcggtcaca acctcgtgga 180
cgccgacaac ctatccccaa ggctcgccgg cccgaggga gggcctgggc ccagcccggg 240
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cgtggctccc ggcctagttg gggcccccacg gaccccccggc gtaggtcgcg caatttgggt 360
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ctgctgtcct gttgaccat cccagcttcc gct 573

<210> 111
<211> 573
<212> DNA
<213> Homo sapiens

<220>
<223> Individual Isolate: D1

<400> 111
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gacgtcaagt tcccggcg gggtcagatc gttgggtggag tttacctgtt gccgcgcagg 120
ggccccaggt tgggtgtgcg cgcgactagg aagacttccg agcggtcaca acctcgtgga 180
aggcgacaac ctatccccaa ggctcgccgg cccgagggtt gggcctgggc tcagcccggg 240
tacccttggc ccctctatgg caacgagggc ttgggtggg caggtggct cctgtcaccc 300
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ggcgccccc taggggtgc tgccagggcc ctggcgcatg gcgtccgggt tctggaggac 480
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ttgctgtcct gttgaccat cccagcttcc gct 573

<210> 112
<211> 573
<212> DNA
<213> Homo sapiens

<220>
<223> Individual Isolate: US6

<400> 112
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gacgtcaagt tcccggcg gggtcagatc gttgggtggag tttacctgtt gccgcgcagg 120
ggccccaggt tgggtgtgcg cgcgactagg aagacttccg agcggtcaca acctcgtgga 180
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ggcgtgaact atgcaacagg gaacttgccc gtttgcctt tctctatctt cctcttggct 540
ttgctgtcct gtttgcaccat tccagcttcc gct 573

<210> 113
<211> 573
<212> DNA
<213> Homo sapiens

<220>
<223> Individual Isolate: P10

<400> 113
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gacgtcaagt tcccgggcgg tggtcagatc gttgggtggag tttacctgtt gccgcgcagg 120
ggccccaggt tgggtgtgcg cgcgactagg aagacttccg agcggtcgca acctcgtgga 180
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ttgctgtcct ggcctgaccat cccagcggtcc gct 573

<210> 114
<211> 573
<212> DNA
<213> Homo sapiens

<220>
<223> Individual Isolate: DK1

<400> 114
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gacgtcaagt tcccgggcgg tggtcagatc gttgggtggag tttacctgtt gccgcgcagg 120
ggccccaggt tgggtgtgcg cgcgactagg aagacttccg agcggtcgca acctcgtgga 180
aggcgacaac ctatccccaa ggctcgccgg cccgagggca gggcctgggc tcagccccc 240
tacccttggc ccctctatgg caatgagggc atggggtggg caggatggct cctgtcaccc 300
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ggcgtgaact acgcaacagg gaatttgccc gtttgcctt tctctatctt cctcttggct 540
ctgttgcctt gtttgcaccat cccagcttcc gcc 573

<210> 115
<211> 573
<212> DNA
<213> Homo sapiens

<220>
<223> Individual Isolate: T10

<400> 115
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cgtggctccc ggcctagttg gggccccaca gaccccccggc gtaggtcgcg taatttgggt 360
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ggcgtgaact atgcaacagg gaatttgccc ggttgctctt tttctatctt cctcttggct 540
ctgctgtctt gtctgaccat cccagcttcc gct 573

<210> 116
<211> 573
<212> DNA
<213> Homo sapiens

<220>
<223> Individual Isolate: SW2

<400> 116
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gacgtcaagt tcccgggcgg tggccagatc gttggtggag tttacctgtt gccgcgcagg 120
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ttgctgtccct gtctgaccat cccagcttcc gct 573

<210> 117
<211> 573
<212> DNA
<213> Homo sapiens

<220>

<223> Individual Isolate: IND3

<400> 117

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gacgtcaagt tcccggcgg tggccagatc gttggtggag tttacctgtt gccgcgcagg 120
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ttgctatcct gttgaccat cccagcttcc gct 573

<210> 118

<211> 573

<212> DNA

<213> Homo sapiens

<220>

<223> Individual Isolate: IND8

<400> 118

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gacgtcaagt tcccggcgg tggccagatc gttggtggag tttacctgtt gccgcgcagg 120
ggccccaggt tgggtgtgcg cgcgactagg aagacttccg agcggtcgca acctcgtgga 180
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ttgctatcct gttgaccgt cccagcttcc gct 573

<210> 119

<211> 573

<212> DNA

<213> Homo sapiens

<220>

<223> Individual Isolate: S9

<400> 119

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gacgttaagt tcccgggcgg tggtcagatc gtcggtggag tttacctgtt gccgcgcagg 120
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ttgctgtcct gtttgaccat cccagttcc gct 573

<210> 120

<211> 573

<212> DNA

<213> Homo sapiens

<220>

<223> Individual Isolate: HK3

<400> 120

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ggccccaggt tgggtgtgcg cgcgaccagg aagacttcag agcggtcgca acctcgtgga 180
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ggtgccccc tagggggcgt tgccagagcc ttggcacatg gtgtccgggt tctggaggac 480
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ttgctgtcct gtttgaccac cccagttcc gct 573

<210> 121

<211> 573

<212> DNA

<213> Homo sapiens

<220>

<223> Individual Isolate: HK5

<400> 121

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ggcgccccc tagggggcgt tgccagagcc ctggcacacg gtgtccgggt tctggaggac 480
ggcgtgaact acgcaacagg gaatataccc ggttgctctt tctctatctt cctttggct 540
ttgctgtcct gtctgaccac cccagttcc gct 573

<210> 122
<211> 573
<212> DNA
<213> Homo sapiens

<220>
<223> Individual Isolate: HK4

<400> 122
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gacgttaagt tcccgggcgg tggccagatc gtcggtgag tttacctgtt gccgcgcagg 120
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<210> 123
<211> 573
<212> DNA
<213> Homo sapiens

<220>
<223> Individual Isolate: P8

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ggcgccccc tagggggcgt tgccagggcc ctggcgcatg gcttccgggt tggaggac 480
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ttgctgtcct gtctgaccat cccagttcc gct 573

<210> 124

<211> 573
<212> DNA
<213> Homo sapiens

<220>
<223> Individual Isolate: T3

<400> 124
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ggccccaggt tgggtgtgcg cgcgactagg aagacttccg agcggtcgca acctcggtga 180
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cgcggtccc ggcctaattt gggcccccaca gaccccccggc gtaggtcgcg taatctgggt 360
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ttgctgtcct gcttgaccat cccagcttcc gct 573

<210> 125
<211> 573
<212> DNA
<213> Homo sapiens

<220>
<223> Individual Isolate: T4

<400> 125
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gacgttaagt tccccggcgg cggccagatc gttggcggag tatacttgcgtt gccgcgcagg 120
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cgagggtccc gtccctcctg gggccccaat gaccccccggc ataggtcgcg caacgtgggt 360
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ggcgccccgt tgggtggcgt cgccagagct ctcgcgcattg gcgtgagagt cctggaggac 480
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ctactgtcct gcatcaccat tccagtcctcc gct 573

<210> 126
<211> 573
<212> DNA
<213> Homo sapiens

<220>
<223> Individual Isolate: US10

<400> 126

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gacgttaagt ttccgggcgg cggccagatc gttggcggag tataacttgtt gccgcgcagg 120
ggcccccaggt tgggtgtgcg cgcgacaagg aagacttcgg agcggtccca gccacgtggg 180
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tacccttggc ccctatatgg gaatgagggg ctcggctggg cagggatggct cctgtcccc 300
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ggcgctccgc ttggtggcgt cgccagagct ctcgcgcatg gcgtgagggt cctggaggac 480
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ttactgtcct gcatcaccat tccagtctct gct 573

<210> 127

<211> 573

<212> DNA

<213> Homo sapiens

<220>

<223> Individual Isolate: T9

<400> 127

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ggcccttaggt tgggtgtgcg cacgacaagg aagacttcgg agcggtccca gccacgtggg 180
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tacccttggc ctctatatgg gaatgagggg ctcggctggg cgggatggct cctgtcccc 300
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ggcgccccgc ttggtggcgt tgccagagct ctcgcgcacg gcgtgagagt cctggaggac 480
ggggtaatt atgcaacagg gaacctaccc ggttgcctt tttctatctt cttgctggcc 540
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<210> 128

<211> 573

<212> DNA

<213> Homo sapiens

<220>

<223> Individual Isolate: T2

<400> 128

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ggcccccaggt tgggtgtgcg cgcgacaagg aagacttcgg agcggtccca gcctcgtgga 180
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cgaggttctc gtccctcttg gggcccaat gaccccccgc ataggtcgcg caatgtgggt 360
aaagtcatcg ataccctaac gtgcggctt gccgacctca tgggttacat ccccgctcgta 420
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ggagttaatt atgcaacagg taacttaccc ggttgctcct tttctatctt cttgctagcc 540
ctgctgtcct gcatcaactat tccggtttca gct 573

<210> 129
<211> 573
<212> DNA
<213> Homo sapiens

<220>
<223> Individual Isolate: T8

<400> 129
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cttctgtcat gcttcacagt gccagtgtct gca 573

<210> 130
<211> 573
<212> DNA
<213> Homo sapiens

<220>
<223> Individual Isolate: US1

<400> 130
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cttctgtcgt gcgccacggt gccgggtgtct gca 573

<210> 131
<211> 573
<212> DNA
<213> Homo sapiens

<220>
<223> Individual Isolate: DK11

<400> 131
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gacgttaagt tcccgggtgg cggccagatc gttggcggag tttacttgct gccgcgcagg 120
ggccccaggt tgggtgtgcg cacgacaagg aagacttccg agcgatccca gccgcgtggg 180
agacgccagc ccatcccgaa agatcggcgc tccaccggca agccctgggg aaagccagga 240
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cgcgggtctc atcctaattg gggcccccact gaccccccgc ataaatcacg caatttgggt 360
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ggcgccccgg tcggaggcgt cgccagagct ctggcacacg gtgttagagt cctggaagac 480
gggataaaatt acgcaacagg gaatctgcct ggttgctctt tttctatctt cttacttgct 540
cttctgtcat gctgcacagt gccagtgtct gcg 573

<210> 132
<211> 573
<212> DNA
<213> Homo sapiens

<220>
<223> Individual Isolate: SW3

<400> 132
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gacgttaagt tcccgggtgg cggccagatc gttggcggag tttacttgct gccgcgcagg 120
ggccccaggt tgggtgtgcg cgcgacaagg aagacttccg agcgatccca gccgcgtggg 180
agacgccagc ccatcccgaa agatcggcgc tccaccggca agtcctgggg aaagccagga 240
tatccttggc ccctgtatgg aaacgagggc tgcggctggg caggttggct cctgtccccc 300
cgcgggtctc atcctaattg gggcccccact gaccccccgc atagatcacg caatttgggc 360
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ggcgccccgg tcggaggcgt cgccagagct ctggcacacg gtgttagagt cctggaagac 480
gggataaaatt acgcaacagg gaatctgcct ggttgctctt tttctatctt cttacttgct 540
cttctgtcgt gcttcacagt gccagtgtct gcg 573

<210> 133
<211> 573
<212> DNA
<213> Homo sapiens

<220>

<223> Individual Isolate: DK8

<400> 133

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ggcccccaggt tgggtgtgcg cgcgacaagg aagtcttccg agcgatccca gccgcgtggg 180
aggcgccagc ccacccgaa agatcggcgc tccacccggca agtcctgggg aaaaccggga 240
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ggcgccccgg ttggaggcggt cgccagagct ctggcacacg gtgttagggt cctggaagac 480
gggataaaatt acgcaacagg gaatttgctt gggtgctt tttctatctt cttgcttgct 540
cttctgtcggt gctgacagt gccagtgtct gcg 573

<210> 134

<211> 573

<212> DNA

<213> Homo sapiens

<220>

<223> Individual Isolate: S83

<400> 134

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gacgtcaagt tcccgggcgg tggccagatc gttggcggag tataacttgct gccgcgcagg 120
ggcccgagat tgggtgtgcg cgcgacgagg aaaacttccg aacggtccca gccacgtggg 180
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ggcgctcccg ttggcggcggt tgccagagcc ctcggccatg gggtagggt tctggaggac 480
gggataaaatt atgcaacggg gaatttgccc gggtgctt tctctatctt tctcttggcc 540
cttctgtctt gcatctctgt gccagtttcc gcc 573

<210> 135

<211> 573

<212> DNA

<213> Homo sapiens

<220>

<223> Individual Isolate: HK10

<400> 135

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gacgttaagt tcccgggtgg cggacagatc gttggtgag tatacgtgtt gccgcgcagg 120
ggcccacat tgggtgtgcg cgacgcgcgt aaaacttctg aacggtcaca gcctcgcgga 180
cgacgcacgc ctatccccaa ggccgcgtcg agcgaaggcc ggtcctggc tcagccccgg 240
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<210> 136

<211> 573

<212> DNA

<213> Homo sapiens

<220>

<223> Individual Isolate: S52

<400> 136

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ctgttctctt gcttagttca tccgcagct agt 573

<210> 137

<211> 573

<212> DNA

<213> Homo sapiens

<220>

<223> Individual Isolate: S2

<400> 137

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ggcgctcccg taggaggcgt cgcaagagcc ctcgcgcatg gcgtgagggc cttgaagac 480
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ctgttctctt gcttaattca tccagcagct agt 573

<210> 138
<211> 573
<212> DNA
<213> Homo sapiens

<220>
<223> Individual Isolate: DK12

<400> 138
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<210> 139
<211> 573
<212> DNA
<213> Homo sapiens

<220>
<223> Individual Isolate: Z4

<400> 139
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<210> 140

<211> 573
<212> DNA
<213> Homo sapiens

<220>
<223> Individual Isolate: Z8

<400> 140
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cttctctcggt gcctaaccgt cccagcgctt gct 573

<210> 141
<211> 573
<212> DNA
<213> Homo sapiens

<220>
<223> Individual Isolate: Z1

<400> 141
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<210> 142
<211> 573
<212> DNA
<213> Homo sapiens

<220>
<223> Individual Isolate: Z5

<400> 142

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ggaatcaact atgcaacagg gaatcttcctt ggttgcctt tttctatctt cctacttgca 540
ctttctcgt gcttgcacac accggcatcc gct 573

<210> 143

<211> 573

<212> DNA

<213> Homo sapiens

<220>

<223> Individual Isolate: Z6

<400> 143

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gacgttaagt tcccgggtgg tggcagatc gttggcggag tttacttgtt gccgcgcagg 120
ggccccaggt tgggtgtgcg cgcgactcgg aagacttcgg agcggtcgca acctcgccgc 180
agacgcccagc ctatccccaa ggcacgtcga tctgagggaa ggtcctgggc tcagccccgg 240
tatccatggc ctcttacgg taatgagggt tgggtgggg cggatggct cctgtcaccc 300
cgtggctctc gaccgtcttggcccaaaat gatccccggc gaaggtcccg caacttgggt 360
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ggcgccccccg tggcgccgt cgccagggcc ctggcacatg gtgttagggc tggaggac 480
ggatcaatt atgcaacagg gaatcttcctt ggttgcctt tctctatctt cctcttggca 540
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<210> 144

<211> 573

<212> DNA

<213> Homo sapiens

<220>

<223> Individual Isolate: Z7

<400> 144

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gacgttaagt tcccggcggtgg tggcagatc gttggcggag tttacttgtt gccgcgcagg 120
ggccccagat tgggtgtgcg cacaactcgg aagacttcgg agcggtcgca acctcgccgc 180
agacgtcagc ctatccccaa ggcacgtcga tctgagggaa ggtcctgggc tcaacccgg 240

tacccatggc ctcttacgg taacgagggt tgcgggtggg caggtggct cttgtcaccc 300
cgtggctctc gaccgtctt gggccaaat gatccccggc gaaggtcccg caacttgggt 360
aaggtcatcg ataccctaac ctgcggctt gccgaccta tggatacat cccgctcgta 420
ggcgcccccg tgggcggcgt cgccagggcc ctgcgcata gctgttagggc tctggaggac 480
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cttcttcgt gcctgactgt tcccgctcg gcc 573

<210> 145
<211> 573
<212> DNA
<213> Homo sapiens

<220>
<223> Individual Isolate: DK13

<400> 145
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ggcccttagat tgggtgtgcg cgcgactagg aagacttcgg agcggtcgca acctcggtgg 180
aggcgccagc ctatccccaa ggccgcggccaa ctgcgggtta ggtcctggc tcagcctggg 240
tacccttggc cccttacgg caatgagggc tgcgggtggg cgggatggct cctgtcaccc 300
cgtggctctc ggccgtctt gggcccgaaat gatccccggc ggagggtccc caacttgggt 360
aaggtcatcg ataccctaac ttgcggcttc gccgaccta tggatacat cccgctcgta 420
ggcgcccccg tgggtggcgt cgccagagcc ctggcgcata gctgtcaggct tctggaggac 480
ggggtcaatt atgcaacagg gaatcttccc gttgtctt tctctatctt cctcttggca 540
ctgcttcgt gcctgactgt tcccgcttcg gcc 573

<210> 146
<211> 573
<212> DNA
<213> Homo sapiens

<220>
<223> Individual Isolate: SA4

<400> 146
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gacgttaagt tcccggtgg tggtcagatc gttggcggag tctacttgg tccgcgcagg 120
ggcccttaggt tgggtgtgcg cgcgacttcgg aagacttcgg aacggtcgca accccgtggg 180
cgccgcggc ctatccccaa ggccgcggccaa cccacggggc ggtcctgggg tcaacccggg 240
tacccttggc cccttacgc caatgagggc ctgcgggtggg cagggtgttt gctctcccc 300
cgaggctctc ggcctaattt gggcccaat gaccccccggc gaaagtgcgc caatttgggt 360
aaggtcatcg ataccctaac gtgcggattc gccgaccta tgggtacat cccgctcgta 420
ggcgcccccg ttggggcgt cgcaaggggcc cttgcacatg gtgtgagggt tcttgaggac 480
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cttcttcgt gcctgaccgt cccggcctct gca 573

<210> 147
<211> 573
<212> DNA
<213> Homo sapiens

<220>
<223> Individual Isolate: SA5

<400> 147
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gacgtcaagt tcccgggcgg tggtcagatc gttggtggag tttacttgtt gccgcgcagg 120
ggcccttagat tgggtgtgcg cgcgactcgg aagacttcag aacggtcgca accccgtggg 180
cgcgcccagc ctattccaa ggccgcggccaa cccacggggc ggtcctgggg tcaacccggg 240
tacccttggc cccttacgc caatgagggc ctcgggtggg cagggtggtt gctctccccc 300
cgaggctctc ggcctaattg gggcccaat gaccccccggc gaaaatcgcg caatttgggt 360
aaggcatcg ataccctaac gtgcggattc gccgacctca tgggtacat cccgctcgta 420
ggcgcccccg ttggggcgt cgcaagggcc ctcgcacatg gtgtgaggggt tcttgaggac 480
gggttaaact atgcaacagg gaatttgccc gtttgctctt tctctatctt tattttttttt 540
cttctctcgt gcttgaccgt cccagcctct gca 573

<210> 148
<211> 573
<212> DNA
<213> Homo sapiens

<220>
<223> Individual Isolate: SA7

<400> 148
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ggcccttaggt tgggtgtgcg cgcgactcgg aagacttcag aacggtcgca accccgtggg 180
cgcgcccagc ctattccaa ggccgcggccaa cccacggggc ggtcctgggg tcaacccggg 240
tacccttggc cccttacgc caatgagggc ctcgggtggg cagggtggtt gctctccccc 300
cgaggctctc ggcctaattg gggcccaat gaccccccggc gaaagtgcgc caatttgggt 360
aaggcatcg acaccctaac atgcggattc gccgacctca tgggtacat cccgctcgta 420
ggcgcccccg ttggggcgt cgcaagggct ctcgcacacg gtgtgaggggt tcttgaggac 480
gggttaaatt acgcaacagg gaatctgccc gtttgctctt tctctatctt tattttttttt 540
cttctctcgt gcttgaccgt cccagcctcc gca 573

<210> 149
<211> 573
<212> DNA
<213> Homo sapiens

<220>

<223> Individual Isolate: SA1

<400> 149

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gacgtcaagt tcccggcgg tggtcagatc gttggtggag tttacttgtt gccgcgcagg 120
ggccccaggt tgggtgtgcg cgcgactcgg aagacttcgg aacggtcga accccgtggg 180
cggcgccagc ctattccaa ggcgcccaa cccacggcc ggtcctgggg tcaacccggg 240
tacccttggc cccttacgc caatgagggc ctcgggtggg cagggtggtt gctctcccc 300
cgaggctctc ggcctaattg gggcccaat gaccccccggc ggaagtcgcg caatttgggt 360
aaggtcatcg ataccctaac gtgcggattc gccgaccta tgggtacat cccgctcgta 420
ggcgcccccg ttggggcgt cgcaagggtctcgacacg gtgtgagggt tcttgaggac 480
gggttaaact acgcaacagg gaatttgccc gttgctt tctctatctt tattccttgca 540
cttcttcct gtctgatcat cccggcctct gca 573

<210> 150

<211> 573

<212> DNA

<213> Homo sapiens

<220>

<223> Individual Isolate: SA3

<400> 150

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gacgtcaagt tcccggcgg tggtcagatc gttggtggag tttacttgtt gccgcgcagg 120
ggccccaggt tgggtgtgcg cgcgactcgg aagacttcag aacggtcga accccgtgga 180
cggcgccagc ctattccaa ggctcgccag cccacggcc ggtcctgggg tcaacccggg 240
tacccttggc cccttacgc caatgagggc ctcgagtggg cagggtggtt gctctcccc 300
cgaggctctc ggcctagttg gggcccaac gaccccccggc gaaatcgcg caatttgggt 360
aaggtcatcg ataccctaac gtgcggattc gccgatcta tgggtacat cccgctcgta 420
ggcgcccccg ttggggcgt cgcaagggtctcgacatg gtgtgagggt tcttgaggac 480
gggttaaact acgcaacagg gaatttaccc gttgctt tctctatctt tattccttgca 540
cttcttcat gcctgaccgt cccggcctct gca 573

<210> 151

<211> 573

<212> DNA

<213> Homo sapiens

<220>

<223> Individual Isolate: SA13

<400> 151

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ggccctaggt tgggtgtgcg cgcaactcgg aagacttcag aacggtcgca accccgtgga 180
cggcgtcagc ctatcccaa ggccgcggcag cccacgggc ggtcctgggg tcaacccggg 240
tacccttggc cccttatgc caatgagggc ctcgggtggg cagggtggtt gctctcccc 300
cgaggctctc ggcctaattt gggcccaat gaccccccggc gaaatcgcg caacttgggt 360
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ggcgcccccg ttggggcgt cgcaagggtt ctcgcacacg gtgtgagggc ctttgaggac 480
gggttaaact atgcaacagg gaatttaccc ggttgctt tcttatctt tatttttgca 540
cttcttcat gcctgactgt cccgacctct gca 573

<210> 152

<211> 573

<212> DNA

<213> Homo sapiens

<220>

<223> Individual Isolate: SA6

<400> 152

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gacgtcaagt tcccgggcgg tggtcagatc gttggtggag tttacttgtt gccgcgcagg 120
ggccctcgta tgggtgtgcg cgcaactcgg aagacttcag aacggtcgca accccgtgga 180
cggcgtcagc ctatcccaa ggccgcggcag cccacgggc ggtcctgggg tcaacccggg 240
tacccttggc cccttatgc caatgagggc ctcgggtggg cagggtggtt gctctcccc 300
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ggcgcccccg ttggggcgt cgcaagggtt ctcgcacacg gtgtgagggc ctttgaggac 480
gggttaaact atgcaacagg gaatttgcgg ggttgctt tcttatctt tatttttgca 540
cttctctcggt gcctaaccgt ccctgcctt gca 573

<210> 153

<211> 573

<212> DNA

<213> Homo sapiens

<220>

<223> Individual Isolate: SA11

<400> 153

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ggccctaggt tgggtgtgcg cgcaactcgg aagacttcag aacggtcgca accccgtgga 180
cggcgtcagc ctatcccaa ggccgcggcag cccacgggc ggtcctgggg tcaacccggg 240
tacccttggc cccttatgc caatgagggc ctcgggtggg cagggtggtt gctctcccc 300
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ggcggccccc ttggggcggt cgcaagggcc ctcgcacacg gtgtgagagc tcttgaggac 480
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cttctctcggt gcttgaccgt cccggccact gca 573

<210> 154
<211> 573
<212> DNA
<213> Homo sapiens

<220>
<223> Individual Isolate: HK2

<400> 154
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ggcccccgggt tgggtgtgcg cgcgacgaga aagacttccg agcgatccca gcccagaggc 180
aggcgccaaac ctataccaa ggcgcgcccag ccccaggca ggcactggc tcagcccgaa 240
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cgcgctccc ggccacattt gggccccaat gaccccccgc gtcgatcccg gaatttgggt 360
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ggcgccctt tggggccgggt cgcggtgcg ctcgcacatg gcgtgaggc aatcgaggac 480
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ctactctcggt gcctcacaac gccagcttcg gct 573

<210> 155
<211> 191
<212> PRT
<213> Homo sapiens

<220>
<223> Individual Isolate: DK7

<400> 155
Met Ser Thr Asn Pro Lys Pro Gln Arg Lys Thr Lys Arg Asn Thr Asn
1 5 10 15

Arg Arg Pro Gln Asp Val Lys Phe Pro Gly Gly Gly Gln Ile Val Gly
20 25 30

Gly Val Tyr Leu Leu Pro Arg Arg Gly Pro Arg Leu Gly Val Arg Ala
35 40 45

Pro Arg Lys Thr Ser Glu Arg Ser Gln Pro Arg Gly Arg Arg Gln Pro
50 55 60

Ile Pro Lys Ala Arg Arg Pro Glu Gly Arg Thr Trp Ala Gln Pro Gly

65 70 75 80

Tyr Pro Trp Pro Leu Tyr Gly Asn Glu Gly Cys Gly Trp Ala Gly Trp
85 90 95

Leu Leu Ser Pro Arg Gly Ser Arg Pro Ser Trp Gly Pro Thr Asp Pro
100 105 110

Arg Arg Arg Ser Arg Asn Leu Gly Lys Val Ile Asp Thr Leu Thr Cys
115 120 125

Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Ala Pro Leu
130 135 140

Gly Gly Ala Ala Arg Ala Leu Ala His Gly Val Arg Val Leu Glu Asp
145 150 155 160

Gly Val Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe Ser Ile
165 170 175

Phe Leu Leu Ala Leu Leu Ser Cys Leu Thr Val Pro Ala Ser Ala
180 185 190

<210> 156

<211> 191

<212> PRT

<213> Homo sapiens

<220>

<223> Individual Isolate: US11

<400> 156

Met Ser Thr Asn Pro Lys Pro Gln Arg Lys Thr Lys Arg Asn Thr Asn
1 5 10 15

Arg Arg Pro Gln Asp Val Lys Phe Pro Gly Gly Gln Ile Val Gly
20 25 30

Gly Val Tyr Leu Leu Pro Arg Arg Gly Pro Arg Leu Gly Val Arg Ala
35 40 45

Thr Arg Lys Thr Ser Glu Arg Ser Gln Pro Arg Gly Arg Arg Gln Pro
50 55 60

Ile Pro Lys Ala Arg Arg Pro Glu Gly Arg Thr Trp Ala Gln Pro Gly
65 70 75 80

Tyr Pro Trp Pro Leu Tyr Gly Asn Glu Gly Cys Gly Trp Ala Gly Trp
85 90 95

Leu Leu Ser Pro Arg Gly Ser Arg Pro Ser Trp Gly Pro Thr Asp Pro
100 105 110

Arg Arg Arg Ser Arg Asn Leu Gly Lys Val Ile Asp Thr Leu Thr Cys
115 120 125

Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Ala Pro Leu
130 135 140

Gly Gly Ala Ala Arg Ala Leu Ala His Gly Val Arg Val Leu Glu Asp
145 150 155 160

Gly Val Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe Ser Ile
165 170 175

Phe Leu Leu Ala Leu Leu Ser Cys Leu Thr Val Pro Ala Ser Ala
180 185 190

<210> 157

<211> 191

<212> PRT

<213> Homo sapiens

<220>

<223> Individual Isolate: S14

<400> 157

Met Ser Thr Asn Pro Lys Pro Gln Arg Lys Thr Lys Arg Asn Thr Asn
1 5 10 15

Arg Arg Pro Gln Asp Val Lys Phe Pro Gly Gly Gln Ile Val Gly
20 25 30

Gly Val Tyr Leu Leu Pro Arg Arg Gly Pro Arg Leu Gly Val Arg Ala
35 40 45

Thr Arg Lys Thr Ser Glu Arg Ser Gln Pro Arg Gly Arg Arg Gln Pro
50 55 60

Ile Pro Lys Ala Arg Arg Pro Glu Gly Arg Thr Trp Ala Gln Pro Gly
65 70 75 80

Tyr Pro Trp Pro Leu Tyr Gly Asn Glu Gly Cys Gly Trp Ala Gly Trp
85 90 95

Leu Leu Ser Pro Arg Gly Ser Arg Pro Ser Trp Gly Pro Thr Asp Pro
100 105 110

Arg Arg Arg Ser Arg Asn Leu Gly Lys Val Ile Asp Thr Leu Thr Cys
115 120 125

Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Ala Pro Leu
130 135 140

Gly Gly Ala Ala Arg Ala Leu Ala His Gly Val Arg Val Leu Glu Asp
145 150 155 160

Gly Val Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe Ser Ile
165 170 175

Phe Leu Leu Ala Leu Leu Ser Cys Leu Thr Val Pro Ala Ser Ala
180 185 190

<210> 158

<211> 191

<212> PRT

<213> Homo sapiens

<220>

<223> Individual Isolate: SW1

<400> 158

Met Ser Thr Asn Pro Lys Pro Gln Arg Lys Thr Lys Arg Asn Thr Asn
1 5 10 15

Arg Arg Pro Gln Asp Val Lys Phe Pro Gly Gly Gln Ile Val Gly
20 25 30

Gly Val Tyr Leu Leu Pro Arg Arg Gly Pro Arg Leu Gly Val Arg Ala
35 40 45

Thr Arg Lys Thr Ser Glu Arg Ser Gln Pro Arg Gly Arg Arg Gln Pro
50 55 60

Ile Pro Lys Ala Arg Arg Pro Glu Gly Arg Thr Trp Ala Gln Pro Gly
65 70 75 80

Tyr Pro Trp Pro Leu Tyr Gly Asn Glu Gly Cys Gly Trp Ala Gly Trp
85 90 95

Leu Leu Ser Pro Arg Gly Ser Arg Pro Ser Trp Gly Pro Thr Asp Pro

100 105 110

Arg Arg Arg Ser Arg Asn Leu Gly Lys Val Ile Asp Thr Leu Thr Cys
115 120 125

Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Ala Pro Leu
130 135 140

Gly Gly Ala Ala Arg Ala Leu Ala His Gly Val Arg Val Leu Glu Asp
145 150 155 160

Gly Val Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe Ser Ile
165 170 175

Phe Leu Leu Ala Leu Leu Ser Cys Leu Thr Val Pro Ala Ser Ala
180 185 190

<210> 159

<211> 191

<212> PRT

<213> Homo sapiens

<220>

<223> Individual Isolate: S18

<400> 159

Met Ser Thr Asn Pro Lys Pro Gln Arg Lys Thr Lys Arg Asn Thr Asn
1 5 10 15

Arg Arg Pro Gln Asp Val Lys Phe Pro Gly Gly Gln Ile Val Gly
20 25 30

Gly Val Tyr Leu Leu Pro Arg Arg Gly Pro Arg Leu Gly Val Arg Ala
35 40 45

Thr Arg Lys Thr Ser Glu Arg Ser Gln Pro Arg Gly Arg Arg Gln Pro
50 55 60

Ile Pro Lys Ala Arg Arg Pro Glu Gly Arg Thr Trp Ala Gln Pro Gly
65 70 75 80

Tyr Pro Trp Pro Leu Tyr Gly Asn Glu Gly Cys Gly Trp Ala Gly Trp
85 90 95

Leu Leu Ser Pro Arg Gly Ser Arg Pro Ser Trp Gly Pro Thr Asp Pro
100 105 110

Arg Arg Arg Ser Arg Asn Leu Gly Lys Val Ile Asp Thr Leu Thr Cys
115 120 125

Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Ala Pro Leu
130 135 140

Gly Gly Ala Ala Arg Ala Leu Ala His Gly Val Arg Val Leu Glu Asp
145 150 155 160

Gly Val Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe Ser Ile
165 170 175

Phe Leu Leu Ala Leu Leu Ser Cys Leu Thr Val Pro Ala Ser Ala
180 185 190

<210> 160

<211> 191

<212> PRT

<213> Homo sapiens

<220>

<223> Individual Isolate: DR4

<400> 160

Met Ser Thr Asn Pro Lys Pro Gln Arg Lys Thr Lys Arg Asn Thr Asn
1 5 10 15

Arg Arg Pro Gln Asp Val Lys Phe Pro Gly Gly Gln Ile Val Gly
20 25 30

Gly Val Tyr Leu Leu Pro Arg Arg Gly Pro Arg Leu Gly Val Arg Ala
35 40 45

Thr Arg Lys Thr Ser Glu Arg Ser Gln Pro Arg Gly Arg Arg Gln Pro
50 55 60

Ile Pro Lys Ala Arg Arg Pro Glu Gly Arg Thr Trp Ala Gln Pro Gly
65 70 75 80

Tyr Pro Trp Pro Leu Tyr Gly Asn Glu Gly Cys Gly Trp Ala Gly Trp
85 90 95

Leu Leu Ser Pro Arg Gly Ser Arg Pro Ser Trp Gly Pro Thr Asp Pro
100 105 110

Arg Arg Arg Ser Arg Asn Leu Gly Lys Val Ile Asp Thr Leu Thr Cys
115 120 125

Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Ala Pro Leu
130 135 140

Gly Gly Ala Ala Arg Ala Leu Ala His Gly Val Arg Val Leu Glu Asp
145 150 155 160

Gly Val Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe Ser Ile
165 170 175

Phe Leu Leu Ala Leu Leu Ser Cys Leu Thr Val Pro Ala Ser Ala
180 185 190

<210> 161

<211> 191

<212> PRT

<213> Homo sapiens

<220>

<223> Individual Isolate: SA10

<400> 161

Met Ser Thr Asn Pro Lys Pro Gln Arg Lys Thr Lys Arg Asn Thr Asn
1 5 10 15

Arg Arg Pro Gln Asp Val Lys Phe Pro Gly Gly Gln Ile Val Gly
20 25 30

Gly Val Tyr Leu Leu Pro Arg Arg Gly Pro Arg Leu Gly Val Arg Ala
35 40 45

Thr Arg Lys Thr Ser Glu Arg Ser Gln Pro Arg Gly Arg Arg Gln Pro
50 55 60

Ile Pro Lys Ala Arg Gln Pro Glu Gly Arg Thr Trp Ala Gln Pro Gly
65 70 75 80

Tyr Pro Trp Pro Leu Tyr Gly Asn Glu Gly Leu Gly Trp Ala Gly Trp
85 90 95

Leu Leu Ser Pro Arg Gly Ser Arg Pro Ser Trp Gly Pro Thr Asp Pro
100 105 110

Arg Arg Arg Ser Arg Asn Leu Gly Lys Val Ile Asp Thr Leu Thr Cys
115 120 125

Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Ala Pro Leu

130

135

140

Gly Gly Ala Ala Arg Ala Leu Ala His Gly Val Arg Val Leu Glu Asp
145 150 155 160

Gly Val Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Pro Phe Ser Ile
165 170 175

Phe Leu Leu Ala Leu Leu Ser Cys Leu Thr Ile Pro Ala Ser Ala
180 185 190

<210> 162

<211> 191

<212> PRT

<213> Homo sapiens

<220>

<223> Individual Isolate: S45

<400> 162

Met Ser Thr Asn Pro Lys Pro Gln Arg Ala Thr Lys Arg Asn Thr Asn
1 5 10 15

Arg Arg Pro Gln Asp Val Lys Phe Pro Gly Gly Gln Ile Val Gly
20 25 30

Gly Val Tyr Leu Leu Pro Arg Arg Gly Pro Arg Leu Gly Val Arg Ala
35 40 45

Thr Arg Lys Thr Ser Glu Arg Ser Gln Pro Arg Gly Arg Arg Gln Pro
50 55 60

Ile Pro Lys Ala Arg Arg Pro Glu Gly Arg Ala Trp Ala Gln Pro Gly
65 70 75 80

His Pro Trp Pro Leu Tyr Gly Asn Glu Gly Leu Gly Trp Ala Gly Trp
85 90 95

Leu Leu Ser Pro Arg Gly Ser Arg Pro Ser Trp Gly Pro Thr Asp Pro
100 105 110

Arg Arg Arg Ser Arg Asn Leu Gly Lys Val Ile Asp Thr Leu Thr Cys
115 120 125

Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Ala Pro Leu
130 135 140

Gly Gly Ala Ala Arg Ala Leu Ala His Gly Val Arg Val Leu Glu Asp
145 150 155 160

Gly Val Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe Ser Ile
165 170 175

Phe Leu Leu Ala Leu Leu Ser Cys Leu Thr Ile Pro Ala Ser Ala
180 185 190

<210> 163

<211> 191

<212> PRT

<213> Homo sapiens

<220>

<223> Individual Isolate: D1

<400> 163

Met Ser Thr Asn Pro Lys Pro Gln Arg Lys Thr Lys Arg Asn Thr Asn
1 5 10 15

Arg Arg Pro Gln Asp Val Lys Phe Pro Gly Gly Gln Ile Val Gly
20 25 30

Gly Val Tyr Leu Leu Pro Arg Arg Gly Pro Arg Leu Gly Val Arg Ala
35 40 45

Thr Arg Lys Thr Ser Glu Arg Ser Gln Pro Arg Gly Arg Arg Gln Pro
50 55 60

Ile Pro Lys Ala Arg Arg Pro Glu Gly Arg Ala Trp Ala Gln Pro Gly
65 70 75 80

Tyr Pro Trp Pro Leu Tyr Gly Asn Glu Gly Leu Gly Trp Ala Gly Trp
85 90 95

Leu Leu Ser Pro Arg Gly Ser Arg Pro Ser Trp Gly Pro Thr Asp Pro
100 105 110

Arg Arg Arg Ser Arg Asn Leu Gly Lys Val Ile Asp Thr Leu Thr Cys
115 120 125

Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Ala Pro Leu
130 135 140

Gly Gly Ala Ala Arg Ala Leu Ala His Gly Val Arg Val Leu Glu Asp
145 150 155 160

Gly Val Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe Ser Ile
165 170 175

Phe Leu Leu Ala Leu Leu Ser Cys Leu Thr Ile Pro Ala Ser Ala
180 185 190

<210> 164

<211> 191

<212> PRT

<213> Homo sapiens

<220>

<223> Individual Isolate: US6

<400> 164

Met Ser Thr Asn Pro Lys Pro Gln Arg Lys Thr Lys Arg Asn Thr Asn
1 5 10 15

Arg Arg Pro Gln Asp Val Lys Phe Pro Gly Gly Gln Ile Val Gly
20 25 30

Gly Val Tyr Leu Leu Pro Arg Arg Gly Pro Arg Leu Gly Val Arg Ala
35 40 45

Thr Arg Lys Thr Ser Glu Arg Ser Gln Pro Arg Gly Arg Arg Gln Pro
50 55 60

Ile Pro Lys Ala Arg Arg Pro Glu Gly Arg Ala Trp Ala Gln Pro Gly
65 70 75 80

Tyr Pro Trp Pro Leu Tyr Gly Asn Glu Gly Met Gly Trp Ala Gly Trp
85 90 95

Leu Leu Ser Pro Arg Gly Ser Arg Pro Ser Trp Gly Pro Thr Asp Pro
100 105 110

Arg Arg Arg Ser Arg Asn Leu Gly Lys Val Ile Asp Thr Leu Thr Cys
115 120 125

Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Ala Pro Leu
130 135 140

Gly Gly Ala Ala Arg Ala Leu Ala His Gly Val Arg Val Leu Glu Asp
145 150 155 160

Gly Val Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe Ser Ile

165

170

175

Phe Leu Leu Ala Leu Leu Ser Cys Leu Thr Ile Pro Ala Ser Ala
180 185 190

<210> 165

<211> 191

<212> PRT

<213> Homo sapiens

<220>

<223> Individual Isolate: P10

<400> 165

Met Ser Thr Asn Pro Lys Pro Gln Arg Lys Thr Lys Arg Asn Thr Asn
1 5 10 15

Arg Arg Pro Gln Asp Val Lys Phe Pro Gly Gly Gly Gln Ile Val Gly
20 25 30

Gly Val Tyr Leu Leu Pro Arg Arg Gly Pro Arg Leu Gly Val Arg Ala
35 40 45

Thr Arg Lys Thr Ser Glu Arg Ser Gln Pro Arg Gly Arg Arg Gln Pro
50 55 60

Ile Pro Lys Ala Arg Arg Pro Glu Gly Arg Ala Trp Ala Gln Pro Gly
65 70 75 80

Tyr Pro Trp Pro Leu Tyr Gly Asn Glu Gly Leu Gly Trp Ala Gly Trp
85 90 95

Leu Leu Ser Pro Arg Gly Ser Arg Pro Ser Trp Gly Pro Thr Asp Pro
100 105 110

Arg Arg Arg Ser Arg Asn Leu Gly Lys Val Ile Asp Thr Leu Thr Cys
115 120 125

Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Ala Pro Leu
130 135 140

Gly Gly Ala Ala Arg Ala Leu Ala His Gly Val Arg Val Leu Glu Asp
145 150 155 160

Gly Val Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe Ser Ile
165 170 175

Phe Leu Leu Ala Leu Leu Ser Cys Leu Thr Ile Pro Ala Ser Ala
180 185 190

<210> 166

<211> 191

<212> PRT

<213> Homo sapiens

<220>

<223> Individual Isolate: DK1

<400> 166

Met Ser Thr Asn Pro Lys Pro Gln Arg Lys Thr Lys Arg Asn Thr Asn
1 5 10 15

Arg Arg Pro Gln Asp Val Lys Phe Pro Gly Gly Gln Ile Val Gly
20 25 30

Gly Val Tyr Leu Leu Pro Arg Arg Gly Pro Arg Leu Gly Val Arg Ala
35 40 45

Thr Arg Lys Thr Ser Glu Arg Ser Gln Pro Arg Gly Arg Arg Gln Pro
50 55 60

Ile Pro Lys Ala Arg Arg Pro Glu Gly Arg Ala Trp Ala Gln Pro Gly
65 70 75 80

Tyr Pro Trp Pro Leu Tyr Gly Asn Glu Gly Met Gly Trp Ala Gly Trp
85 90 95

Leu Leu Ser Pro Arg Gly Ser Arg Pro Ser Trp Gly Pro Asn Asp Pro
100 105 110

Arg Arg Arg Ser Arg Asn Leu Gly Lys Val Ile Asp Thr Leu Thr Cys
115 120 125

Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Ala Pro Leu
130 135 140

Gly Gly Ala Ala Arg Ala Leu Ala His Gly Val Arg Val Leu Glu Asp
145 150 155 160

Gly Val Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe Ser Ile
165 170 175

Phe Leu Leu Ala Leu Leu Ser Cys Leu Thr Ile Pro Ala Ser Ala
180 185 190

<210> 167
<211> 191
<212> PRT
<213> Homo sapiens

<220>
<223> Individual Isolate: T10

<400> 167
Met Ser Thr Asn Pro Lys Pro Gln Arg Lys Thr Lys Arg Asn Thr Asn
1 5 10 15

Arg Arg Pro Gln Asp Val Lys Phe Pro Gly Gly Gln Ile Val Gly
20 25 30

Gly Val Tyr Leu Leu Pro Arg Arg Gly Pro Arg Leu Gly Val Arg Ala
35 40 45

Thr Arg Lys Thr Ser Glu Arg Ser Gln Pro Arg Gly Arg Arg Gln Pro
50 55 60

Ile Pro Lys Ala Arg Gln Pro Glu Gly Arg Ala Trp Ala Gln Pro Gly
65 70 75 80

Tyr Pro Trp Pro Leu Tyr Gly Asn Glu Gly Met Gly Trp Ala Gly Trp
85 90 95

Leu Leu Ser Pro Arg Gly Ser Arg Pro Ser Trp Gly Pro Thr Asp Pro
100 105 110

Arg Arg Arg Ser Arg Asn Leu Gly Lys Val Ile Asp Thr Leu Thr Cys
115 120 125

Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Ala Pro Leu
130 135 140

Gly Gly Ala Ala Arg Ala Leu Ala His Gly Val Arg Val Leu Glu Asp
145 150 155 160

Gly Val Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe Ser Ile
165 170 175

Phe Leu Leu Ala Leu Leu Ser Cys Leu Thr Ile Pro Ala Ser Ala
180 185 190

<210> 168
<211> 191
<212> PRT
<213> Homo sapiens

<220>
<223> Individual Isolate: SW2

<400> 168
Met Ser Thr Asn Pro Lys Pro Gln Arg Lys Thr Lys Arg Asn Thr Asn
1 5 10 15
Arg Arg Pro Gln Asp Val Lys Phe Pro Gly Gly Gln Ile Val Gly
20 25 30
Gly Val Tyr Leu Leu Pro Arg Arg Gly Pro Arg Leu Gly Val Arg Ala
35 40 45
Thr Arg Lys Thr Ser Glu Arg Ser Gln Pro Arg Gly Arg Arg Gln Pro
50 55 60
Ile Pro Lys Ala Arg Gln Pro Glu Gly Arg Ala Trp Ala Gln Pro Gly
65 70 75 80
Tyr Pro Trp Pro Leu Tyr Gly Asn Glu Gly Met Gly Trp Ala Gly Trp
85 90 95
Leu Leu Ser Pro Arg Gly Ser Arg Pro Ser Trp Gly Pro Thr Asp Pro
100 105 110
Arg Arg Arg Ser Arg Asn Leu Gly Lys Val Ile Asp Thr Leu Thr Cys
115 120 125
Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Ala Pro Leu
130 135 140
Gly Gly Ala Ala Arg Ala Leu Ala His Gly Val Arg Val Leu Glu Asp
145 150 155 160
Gly Val Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe Ser Ile
165 170 175
Phe Leu Leu Ala Leu Leu Ser Cys Leu Thr Ile Pro Ala Ser Ala
180 185 190

<210> 169
<211> 191

<212> PRT

<213> Homo sapiens

<220>

<223> Individual Isolate: IND3

<400> 169

Met Ser Thr Asn Pro Lys Pro Gln Arg Lys Thr Lys Arg Asn Thr Asn
1 5 10 15

Arg Arg Pro Gln Asp Val Lys Phe Pro Gly Gly Gln Ile Val Gly
20 25 30

Gly Val Tyr Leu Leu Pro Arg Arg Gly Pro Arg Leu Gly Val Arg Ala
35 40 45

Thr Arg Lys Thr Ser Glu Arg Ser Gln Pro Arg Gly Arg Arg Gln Pro
50 55 60

Ile Pro Lys Ala Arg Arg Pro Glu Gly Arg Ala Trp Ala Gln Pro Gly
65 70 75 80

Tyr Pro Trp Pro Leu Tyr Gly Asn Glu Gly Leu Gly Trp Ala Gly Trp
85 90 95

Leu Leu Ser Pro Arg Gly Ser Arg Pro Ser Trp Gly Pro Thr Asp Pro
100 105 110

Arg Arg Arg Ser Arg Asn Leu Gly Lys Val Ile Asp Thr Leu Thr Cys
115 120 125

Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Ala Pro Leu
130 135 140

Gly Gly Ala Ala Arg Ala Leu Ala His Gly Val Arg Val Leu Glu Asp
145 150 155 160

Gly Val Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe Ser Ile
165 170 175

Phe Leu Leu Ala Leu Leu Ser Cys Leu Thr Ile Pro Ala Ser Ala
180 185 190

<210> 170

<211> 191

<212> PRT

<213> Homo sapiens

<220>

<223> Individual Isolate: IND8

<400> 170

Met Ser Thr Asn Pro Lys Pro Gln Arg Lys Thr Lys Arg Asn Thr Asn
1 5 10 15

Arg Arg Pro Gln Asp Val Lys Phe Pro Gly Gly Gln Ile Val Gly
20 25 30

Gly Val Tyr Leu Leu Pro Arg Arg Gly Pro Arg Leu Gly Val Arg Ala
35 40 45

Thr Arg Lys Thr Ser Glu Arg Ser Gln Pro Arg Gly Arg Arg Gln Pro
50 55 60

Ile Pro Lys Ala Arg Arg Pro Glu Gly Arg Ala Trp Ala Gln Pro Gly
65 70 75 80

His Pro Trp Pro Leu Tyr Gly Asn Glu Gly Leu Gly Trp Ala Gly Trp
85 90 95

Leu Leu Ser Pro Arg Gly Ser Arg Pro Ser Trp Gly Pro Thr Asp Pro
100 105 110

Arg Arg Arg Ser Arg Asn Leu Gly Lys Val Ile Asp Thr Leu Thr Cys
115 120 125

Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Ala Pro Leu
130 135 140

Gly Gly Ala Ala Arg Ala Leu Ala His Gly Val Arg Val Leu Glu Asp
145 150 155 160

Gly Val Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe Ser Ile
165 170 175

Phe Leu Leu Ala Leu Leu Ser Cys Leu Thr Val Pro Ala Ser Ala
180 185 190

<210> 171

<211> 191

<212> PRT

<213> Homo sapiens

<220>

<223> Individual Isolate: S9

<400> 171

Met Ser Thr Asn Pro Lys Pro Gln Arg Lys Thr Lys Arg Asn Thr Asn
1 5 10 15

Arg Arg Pro Gln Asp Val Lys Phe Pro Gly Gly Gln Ile Val Gly
20 25 30

Gly Val Tyr Leu Leu Pro Arg Arg Gly Pro Arg Leu Gly Val Arg Ala
35 40 45

Thr Arg Lys Thr Ser Glu Arg Ser Gln Pro Arg Gly Arg Arg Gln Pro
50 55 60

Ile Pro Lys Ala Arg His Pro Glu Gly Arg Ala Trp Ala Gln Pro Gly
65 70 75 80

Tyr Pro Trp Pro Leu Tyr Gly Asn Glu Gly Leu Gly Trp Ala Gly Trp
85 90 95

Leu Leu Ser Pro Arg Gly Ser Arg Pro Ser Trp Gly Pro Asn Asp Pro
100 105 110

Arg Arg Arg Ser Arg Asn Leu Gly Lys Val Ile Asp Thr Leu Thr Cys
115 120 125

Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Ala Pro Leu
130 135 140

Gly Gly Ala Ala Arg Ala Leu Ala His Gly Val Arg Val Leu Glu Asp
145 150 155 160

Gly Val Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe Ser Ile
165 170 175

Phe Leu Leu Ala Leu Leu Ser Cys Leu Thr Ile Pro Ala Ser Ala
180 185 190

<210> 172

<211> 191

<212> PRT

<213> Homo sapiens

<220>

<223> Individual Isolate: HK3

<400> 172

Met Ser Thr Asn Pro Lys Pro Gln Arg Lys Thr Lys Arg Asn Thr Asn
1 5 10 15

Arg Arg Pro Gln Asp Val Lys Phe Pro Gly Gly Gln Ile Val Gly
20 25 30

Gly Val Tyr Leu Leu Pro Arg Arg Gly Pro Arg Leu Gly Val Arg Ala
35 40 45

Thr Arg Lys Thr Ser Glu Arg Ser Gln Pro Arg Gly Arg Arg Gln Pro
50 55 60

Ile Pro Lys Ala Arg Gln Pro Glu Gly Arg Thr Trp Ala Gln Pro Gly
65 70 75 80

Tyr Pro Trp Pro Leu Tyr Gly Asn Glu Gly Met Gly Trp Ala Gly Trp
85 90 95

Leu Leu Ser Pro Arg Gly Ser Arg Pro Asn Trp Gly Pro Thr Asp Pro
100 105 110

Arg Arg Arg Ser Arg Asn Leu Gly Lys Val Ile Asp Thr Leu Thr Cys
115 120 125

Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Ala Pro Leu
130 135 140

Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Val Leu Glu Asp
145 150 155 160

Gly Val Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe Ser Ile
165 170 175

Phe Leu Leu Ala Leu Leu Ser Cys Leu Thr Thr Pro Ala Ser Ala
180 185 190

<210> 173

<211> 191

<212> PRT

<213> Homo sapiens

<220>

<223> Individual Isolate: HK5

<400> 173

Met Ser Thr Asn Pro Lys Pro Gln Arg Lys Thr Lys Arg Asn Thr Asn

1	5	10	15
Arg Arg Pro Gln Asp Val Lys Phe Pro Gly Gly Gly Gln Ile Val Gly			
20	25	30	
Gly Val Tyr Leu Leu Pro Arg Arg Gly Pro Arg Leu Gly Val Arg Ala			
35	40	45	
Thr Arg Lys Thr Ser Glu Arg Ser Gln Pro Arg Gly Arg Arg Gln Pro			
50	55	60	
Ile Pro Lys Ala Arg Arg Pro Glu Gly Arg Thr Trp Ala Gln Pro Gly			
65	70	75	80
Tyr Pro Trp Pro Leu Tyr Gly Asn Glu Gly Met Gly Trp Ala Gly Trp			
85	90	95	
Leu Leu Ser Pro His Gly Ser Arg Pro Ser Trp Gly Pro Thr Asp Pro			
100	105	110	
Arg Arg Arg Ser Arg Asn Leu Gly Lys Val Ile Asp Thr Leu Thr Cys			
115	120	125	
Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Ala Pro Leu			
130	135	140	
Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Val Leu Glu Asp			
145	150	155	160
Gly Val Asn Tyr Ala Thr Gly Asn Ile Pro Gly Cys Ser Phe Ser Ile			
165	170	175	
Phe Leu Leu Ala Leu Leu Ser Cys Leu Thr Thr Pro Val Ser Ala			
180	185	190	
<210> 174			
<211> 191			
<212> PRT			
<213> Homo sapiens			
<220>			
<223> Individual Isolate: HK4			
<400> 174			
Met Ser Thr Asn Pro Lys Pro Gln Arg Lys Thr Lys Arg Asn Thr Asn			
1	5	10	15

Arg Arg Pro Gln Asp Val Lys Phe Pro Gly Gly Gln Ile Val Gly
20 25 30

Gly Val Tyr Leu Leu Pro Arg Arg Gly Pro Arg Leu Gly Val Arg Ala
35 40 45

Thr Arg Lys Thr Ser Glu Arg Ser Gln Pro Arg Gly Arg Arg Gln Pro
50 55 60

Ile Pro Lys Ala Arg Gln Pro Glu Gly Arg Thr Trp Ala Gln Pro Gly
65 70 75 80

Tyr Pro Trp Pro Leu Tyr Gly Asn Glu Gly Met Gly Trp Ala Gly Trp
85 90 95

Leu Leu Ser Pro Arg Gly Ser Arg Pro Ser Trp Gly Pro Thr Asp Pro
100 105 110

Arg Arg Arg Ser Arg Asn Leu Gly Lys Val Ile Asp Thr Leu Thr Cys
115 120 125

Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Ala Pro Leu
130 135 140

Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Val Val Glu Asp
145 150 155 160

Gly Val Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe Ser Ile
165 170 175

Phe Leu Leu Ala Leu Leu Ser Cys Leu Thr Ile Pro Ala Ser Ala
180 185 190

<210> 175

<211> 191

<212> PRT

<213> Homo sapiens

<220>

<223> Individual Isolate: P8

<400> 175

Met Ser Thr Thr Pro Lys Pro Gln Arg Lys Thr Lys Arg Asn Thr Ser
1 5 10 15

Arg Arg Pro Gln Asp Val Lys Phe Pro Gly Gly Gln Ile Val Gly
20 25 30

Gly Val Tyr Leu Leu Pro Arg Arg Gly Pro Arg Leu Gly Val Arg Ala
35 40 45

Thr Arg Lys Thr Ser Glu Arg Ser Gln Pro Arg Gly Arg Arg Gln Pro
50 55 60

Ile Pro Lys Ala Arg Arg Pro Glu Gly Arg Ala Trp Ala Gln Pro Gly
65 70 75 80

His Pro Trp Pro Leu Tyr Ala Asn Glu Gly Leu Gly Trp Ala Gly Trp
85 90 95

Leu Leu Ser Pro Arg Gly Ser Arg Pro Ser Trp Gly Pro Thr Asp Pro
100 105 110

Arg Arg Arg Ser Arg Asn Leu Gly Lys Val Ile Asp Thr Leu Thr Cys
115 120 125

Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Gly Pro Leu
130 135 140

Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Val Val Glu Asp
145 150 155 160

Gly Val Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe Ser Ile
165 170 175

Phe Leu Leu Ala Leu Leu Ser Cys Leu Thr Ile Pro Ala Ser Ala
180 185 190

<210> 176

<211> 191

<212> PRT

<213> Homo sapiens

<220>

<223> Individual Isolate: T3

<400> 176

Met Ser Thr Asn Pro Lys Pro Gln Arg Lys Thr Lys Arg Asn Thr Asn
1 5 10 15

Arg Arg Pro Gln Asp Val Lys Phe Pro Gly Gly Gln Ile Val Gly
20 25 30

Gly Val Tyr Leu Leu Pro Arg Arg Gly Pro Arg Leu Gly Val Arg Ala

35

40

45

Thr Arg Lys Thr Ser Glu Arg Ser Gln Pro Arg Gly Arg Arg Gln Pro
50 55 60

Ile Pro Lys Ala Arg Arg Pro Glu Gly Arg Ala Trp Ala Gln Pro Gly
65 70 75 80

Tyr Pro Trp Pro Leu Tyr Gly Asp Glu Gly Met Gly Trp Ala Gly Trp
85 90 95

Leu Leu Ser Pro Arg Gly Ser Arg Pro Asn Trp Gly Pro Thr Asp Pro
100 105 110

Arg Arg Arg Ser Arg Asn Leu Gly Lys Val Ile Asp Thr Leu Thr Cys
115 120 125

Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Ala Pro Leu
130 135 140

Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Val Leu Glu Asp
145 150 155 160

Gly Val Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe Ser Ile
165 170 175

Phe Leu Leu Ala Leu Leu Ser Cys Leu Thr Ile Pro Ala Ser Ala
180 185 190

<210> 177

<211> 191

<212> PRT

<213> Homo sapiens

<220>

<223> Individual Isolate: T4

<400> 177

Met Ser Thr Asn Pro Lys Pro Gln Arg Lys Thr Lys Arg Asn Thr Asn
1 5 10 15

Arg Arg Pro Gln Asp Val Lys Phe Pro Gly Gly Gln Ile Val Gly
20 25 30

Gly Val Tyr Leu Leu Pro Arg Arg Gly Pro Arg Leu Gly Val Arg Ala
35 40 45

Thr Arg Lys Thr Ser Glu Arg Ser Gln Pro Arg Gly Arg Arg Gln Pro
50 55 60

Ile Pro Lys Asp Arg Arg Ser Thr Gly Lys Ser Trp Gly Lys Pro Gly
65 70 75 80

Tyr Pro Trp Pro Leu Tyr Gly Asn Glu Gly Leu Gly Trp Ala Gly Trp
85 90 95

Leu Leu Ser Pro Arg Gly Ser Arg Pro Ser Trp Gly Pro Asn Asp Pro
100 105 110

Arg His Arg Ser Arg Asn Val Gly Lys Val Ile Asp Thr Leu Thr Cys
115 120 125

Ser Leu Ala Asp Leu Met Gly Tyr Val Pro Val Val Gly Gly Pro Leu
130 135 140

Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Val Leu Glu Asp
145 150 155 160

Gly Val Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe Ser Ile
165 170 175

Phe Leu Leu Ala Leu Leu Ser Cys Ile Thr Ile Pro Val Ser Ala
180 185 190

<210> 178

<211> 191

<212> PRT

<213> Homo sapiens

<220>

<223> Individual Isolate: US10

<400> 178

Met Ser Thr Asn Pro Lys Pro Gln Arg Lys Thr Lys Arg Asn Thr Asn
1 5 10 15

Arg Arg Pro Gln Asp Val Lys Phe Pro Gly Gly Gln Ile Val Gly
20 25 30

Gly Val Tyr Leu Leu Pro Arg Arg Gly Pro Arg Leu Gly Val Arg Ala
35 40 45

Thr Arg Lys Thr Ser Glu Arg Ser Gln Pro Arg Gly Arg Arg Gln Pro
50 55 60

Ile Pro Lys Asp Arg Arg Pro Thr Gly Lys Ser Trp Gly Lys Pro Gly
65 70 75 80

Tyr Pro Trp Pro Leu Tyr Gly Asn Glu Gly Leu Gly Trp Ala Gly Trp
85 90 95

Leu Leu Ser Pro Arg Gly Ser Arg Pro Ser Trp Gly Pro Thr Asp Pro
100 105 110

Arg His Arg Ser Arg Asn Val Gly Lys Val Ile Asp Thr Leu Thr Cys
115 120 125

Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Val Val Gly Ala Pro Leu
130 135 140

Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Val Leu Glu Asp
145 150 155 160

Gly Val Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe Ser Ile
165 170 175

Phe Leu Leu Ala Leu Leu Ser Cys Ile Thr Ile Pro Val Ser Ala
180 185 190

<210> 179

<211> 191

<212> PRT

<213> Homo sapiens

<220>

<223> Individual Isolate: T9

<400> 179

Met Ser Thr Asn Pro Lys Pro Gln Arg Lys Thr Ile Arg Asn Thr Asn
1 5 10 15

Arg Arg Pro Gln Asp Val Lys Phe Pro Gly Gly Gln Ile Val Gly
20 25 30

Gly Val Tyr Leu Leu Pro Arg Arg Gly Pro Arg Leu Gly Val Arg Thr
35 40 45

Thr Arg Lys Thr Ser Glu Arg Ser Gln Pro Arg Gly Arg Arg Gln Pro
50 55 60

Ile Pro Lys Asp Arg Arg Ser Thr Gly Lys Ser Trp Gly Lys Pro Gly

65 70 75 80

Tyr Pro Trp Pro Leu Tyr Gly Asn Glu Gly Leu Gly Trp Ala Gly Trp
85 90 95

Leu Leu Ser Pro Arg Gly Ser Arg Pro Ser Trp Gly Pro Ser Asp Pro
100 105 110

Arg His Arg Ser Arg Asn Val Gly Lys Val Ile Asp Thr Leu Thr Cys
115 120 125

Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Val Val Gly Ala Pro Leu
130 135 140

Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Val Leu Glu Asp
145 150 155 160

Gly Val Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe Ser Ile
165 170 175

Phe Leu Leu Ala Leu Leu Ser Cys Ile Thr Thr Pro Ala Ser Ala
180 185 190

<210> 180

<211> 191

<212> PRT

<213> Homo sapiens

<220>

<223> Individual Isolate: T2

<400> 180

Met Ser Thr Ile Pro Lys Pro Gln Arg Lys Thr Lys Arg Asn Thr Asn
1 5 10 15

Arg Arg Pro Gln Asp Val Lys Phe Pro Gly Gly Gln Ile Val Gly
20 25 30

Gly Val Tyr Leu Leu Pro Arg Arg Gly Pro Arg Leu Gly Val Arg Ala
35 40 45

Thr Arg Lys Thr Ser Glu Arg Ser Gln Pro Arg Gly Arg Arg Gln Pro
50 55 60

Ile Pro Lys Asp Arg Arg Ser Thr Gly Lys Ser Trp Gly Lys Pro Gly
65 70 75 80

Tyr Pro Trp Pro Leu Tyr Gly Asn Glu Gly Leu Gly Trp Ala Gly Trp
85 90 95

Leu Leu Ser Pro Arg Gly Ser Arg Pro Ser Trp Gly Pro Asn Asp Pro
100 105 110

Arg His Arg Ser Arg Asn Val Gly Lys Val Ile Asp Thr Leu Thr Cys
115 120 125

Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Val Val Gly Ala Pro Leu
130 135 140

Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Val Leu Glu Asp
145 150 155 160

Gly Val Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe Ser Ile
165 170 175

Phe Leu Leu Ala Leu Leu Ser Cys Ile Thr Ile Pro Val Ser Ala
180 185 190

<210> 181

<211> 191

<212> PRT

<213> Homo sapiens

<220>

<223> Individual Isolate: T8

<400> 181

Met Ser Thr Asn Pro Lys Pro Gln Arg Lys Thr Lys Arg Asn Thr Asn
1 5 10 15

Arg Arg Pro Gln Asp Val Lys Phe Pro Gly Gly Gln Ile Val Gly
20 25 30

Gly Val Tyr Leu Leu Pro Arg Arg Gly Pro Arg Leu Gly Val Arg Ala
35 40 45

Thr Arg Lys Thr Ser Glu Arg Ser Gln Pro Arg Gly Arg Arg Gln Pro
50 55 60

Ile Pro Lys Asp Arg Arg Ser Thr Gly Lys Ser Trp Gly Lys Pro Gly
65 70 75 80

Tyr Pro Trp Pro Leu Tyr Gly Asn Glu Gly Cys Gly Trp Ala Gly Trp
85 90 95

Leu Leu Ser Pro Arg Gly Ser Arg Pro Thr Trp Gly Pro Thr Asp Pro
100 105 110

Arg His Arg Ser Arg Asn Leu Gly Arg Val Ile Asp Thr Ile Thr Cys
115 120 125

Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Val Val Gly Ala Pro Val
130 135 140

Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Val Leu Glu Asp
145 150 155 160

Gly Ile Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe Ser Ile
165 170 175

Phe Leu Leu Ala Leu Leu Ser Cys Phe Thr Val Pro Val Ser Ala
180 185 190

<210> 182

<211> 191

<212> PRT

<213> Homo sapiens

<220>

<223> Individual Isolate: US1

<400> 182

Met Ser Thr Asn Pro Lys Pro Gln Arg Lys Thr Lys Arg Asn Thr Asn
1 5 10 15

Arg Arg Pro Gln Asp Val Lys Phe Pro Gly Gly Gln Ile Val Gly
20 25 30

Gly Val Tyr Leu Leu Pro Arg Arg Gly Pro Arg Leu Gly Val Arg Ala
35 40 45

Thr Arg Lys Thr Ser Glu Arg Ser Gln Pro Arg Gly Arg Arg Gln Pro
50 55 60

Ile Pro Lys Asp Arg Arg Ser Thr Gly Lys Ser Trp Gly Lys Pro Gly
65 70 75 80

Tyr Pro Trp Pro Leu Tyr Gly Asn Glu Gly Cys Gly Trp Ala Gly Trp
85 90 95

Leu Leu Ser Pro Arg Gly Ser Arg Pro Thr Trp Gly Pro Thr Asp Pro

100	105	110
Arg His Arg Ser Arg Asn Leu Gly Lys Val Ile Asp Thr Ile Thr Cys		
115	120	125
Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Val Val Gly Ala Pro Val		
130	135	140
Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Val Leu Glu Asp		
145	150	155
Gly Ile Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe Ser Ile		
165	170	175
Phe Leu Leu Ala Leu Leu Ser Cys Ala Thr Val Pro Val Ser Ala		
180	185	190
<210> 183		
<211> 191		
<212> PRT		
<213> Homo sapiens		
<220>		
<223> Individual Isolate: DK11		
<400> 183		
Met Ser Thr Asn Pro Lys Pro Gln Arg Lys Thr Lys Arg Asn Thr Asn		
1	5	10
15		
Arg Arg Pro Gln Asp Val Lys Phe Pro Gly Gly Gln Ile Val Gly		
20	25	30
Gly Val Tyr Leu Leu Pro Arg Arg Gly Pro Arg Leu Gly Val Arg Thr		
35	40	45
Thr Arg Lys Thr Ser Glu Arg Ser Gln Pro Arg Gly Arg Arg Gln Pro		
50	55	60
Ile Pro Lys Asp Arg Arg Ser Thr Gly Lys Pro Trp Gly Lys Pro Gly		
65	70	75
80		
Tyr Pro Trp Pro Leu Tyr Gly Asn Glu Gly Cys Gly Trp Ala Gly Trp		
85	90	95
Leu Leu Ser Pro Arg Gly Ser His Pro Asn Trp Gly Pro Thr Asp Pro		
100	105	110

Arg His Lys Ser Arg Asn Leu Gly Lys Val Ile Asp Thr Ile Thr Cys
115 120 125

Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Val Val Gly Ala Pro Val
130 135 140

Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Val Leu Glu Asp
145 150 155 160

Gly Ile Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe Ser Ile
165 170 175

Phe Leu Leu Ala Leu Leu Ser Cys Cys Thr Val Pro Val Ser Ala
180 185 190

<210> 184

<211> 191

<212> PRT

<213> Homo sapiens

<220>

<223> Individual Isolate: SW3

<400> 184

Met Ser Thr Asn Pro Lys Pro Gln Arg Lys Thr Lys Arg Asn Thr Asn
1 5 10 15

Arg Arg Pro Gln Asp Val Lys Phe Pro Gly Gly Gln Ile Val Gly
20 25 30

Gly Val Tyr Leu Leu Pro Arg Arg Gly Pro Arg Leu Gly Val Arg Ala
35 40 45

Thr Arg Lys Thr Ser Glu Arg Ser Gln Pro Arg Gly Arg Arg Gln Pro
50 55 60

Ile Pro Lys Asp Arg Arg Ser Thr Gly Lys Ser Trp Gly Lys Pro Gly
65 70 75 80

Tyr Pro Trp Pro Leu Tyr Gly Asn Glu Gly Cys Gly Trp Ala Gly Trp
85 90 95

Leu Leu Ser Pro Arg Gly Ser His Pro Asn Trp Gly Pro Thr Asp Pro
100 105 110

Arg His Arg Ser Arg Asn Leu Gly Lys Val Ile Asp Thr Ile Thr Cys
115 120 125

Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Val Val Gly Ala Pro Val
130 135 140

Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Val Leu Glu Asp
145 150 155 160

Gly Ile Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe Ser Ile
165 170 175

Phe Leu Leu Ala Leu Leu Ser Cys Phe Thr Val Pro Val Ser Ala
180 185 190

<210> 185

<211> 191

<212> PRT

<213> Homo sapiens

<220>

<223> Individual Isolate: DK8

<400> 185

Met Ser Thr Asn Pro Lys Pro Gln Arg Lys Thr Lys Arg Asn Thr Asn
1 5 10 15

Arg Arg Pro Gln Asp Val Lys Phe Pro Gly Gly Gln Ile Val Gly
20 25 30

Gly Val Tyr Leu Leu Pro Arg Arg Gly Pro Arg Leu Gly Val Arg Ala
35 40 45

Thr Arg Lys Ser Ser Glu Arg Ser Gln Pro Arg Gly Arg Arg Gln Pro
50 55 60

Ile Pro Lys Asp Arg Arg Ser Thr Gly Lys Ser Trp Gly Lys Pro Gly
65 70 75 80

Tyr Pro Trp Pro Leu Tyr Gly Asn Glu Gly Cys Gly Trp Ala Gly Trp
85 90 95

Leu Leu Ser Pro Arg Gly Ser Arg Pro Thr Trp Gly Pro Thr Asp Pro
100 105 110

Arg His Arg Ser Arg Asn Leu Gly Lys Val Ile Asp Thr Ile Thr Cys
115 120 125

Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Val Val Gly Ala Pro Val

130	135	140
Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Val Leu Glu Asp		
145	150	155
Gly Ile Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe Ser Ile		
165	170	175
Phe Leu Leu Ala Leu Leu Ser Cys Cys Thr Val Pro Val Ser Ala		
180	185	190
<210> 186		
<211> 191		
<212> PRT		
<213> Homo sapiens		
<220>		
<223> Individual Isolate: S83		
<400> 186		
Met Ser Thr Asn Pro Lys Pro Gln Arg Lys Thr Lys Arg Asn Thr Asn		
1	5	10
15		
Arg Arg Pro Gln Asp Val Lys Phe Pro Gly Gly Gln Ile Val Gly		
20	25	30
Gly Val Tyr Leu Leu Pro Arg Arg Gly Pro Arg Leu Gly Val Arg Ala		
35	40	45
Thr Arg Lys Thr Ser Glu Arg Ser Gln Pro Arg Gly Arg Arg Gln Pro		
50	55	60
Ile Pro Lys Asp Arg Arg Thr Thr Gly Lys Ser Trp Gly Arg Pro Gly		
65	70	75
80		
Tyr Pro Trp Pro Leu Tyr Gly Asn Glu Gly Leu Gly Trp Ala Gly Trp		
85	90	95
Leu Leu Ser Pro Arg Gly Ser Arg Pro Ser Trp Gly Pro Thr Asp Pro		
100	105	110
Arg His Lys Ser Arg Asn Leu Gly Lys Val Ile Asp Thr Leu Thr Cys		
115	120	125
Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Val Val Gly Ala Pro Val		
130	135	140

Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Val Leu Glu Asp
145 150 155 160

Gly Ile Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe Ser Ile
165 170 175

Phe Leu Leu Ala Leu Leu Ser Cys Ile Ser Val Pro Val Ser Ala
180 185 190

<210> 187

<211> 191

<212> PRT

<213> Homo sapiens

<220>

<223> Individual Isolate: HK10

<400> 187

Met Ser Thr Leu Pro Lys Pro Gln Arg Lys Thr Lys Arg Asn Thr Ile
1 5 10 15

Arg Arg Pro Gln Asp Val Lys Phe Pro Gly Gly Gln Ile Val Gly
20 25 30

Gly Val Tyr Val Leu Pro Arg Arg Gly Pro Arg Leu Gly Val Arg Ala
35 40 45

Thr Arg Lys Thr Ser Glu Arg Ser Gln Pro Arg Gly Arg Arg Gln Pro
50 55 60

Ile Pro Lys Ala Arg Arg Ser Glu Gly Arg Ser Trp Ala Gln Pro Gly
65 70 75 80

Tyr Pro Trp Pro Leu Tyr Gly Asn Glu Gly Cys Gly Trp Ala Gly Trp
85 90 95

Leu Leu Ser Pro Arg Gly Ser Arg Pro Ser Trp Gly Pro Asn Asp Pro
100 105 110

Arg Arg Arg Ser Arg Asn Leu Gly Lys Val Ile Asp Thr Leu Thr Cys
115 120 125

Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Ala Pro Val
130 135 140

Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Ala Leu Glu Asp
145 150 155 160

Gly Ile Asn Phe Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe Ser Ile
165 170 175

Phe Leu Leu Ala Leu Phe Ser Cys Leu Ile His Pro Ala Ala Ser
180 185 190

<210> 188

<211> 191

<212> PRT

<213> Homo sapiens

<220>

<223> Individual Isolate: S52

<400> 188

Met Ser Thr Leu Pro Lys Pro Gln Arg Lys Thr Lys Arg Asn Thr Ile
1 5 10 15

Arg Arg Pro Gln Asp Val Lys Phe Pro Gly Gly Gln Ile Val Gly
20 25 30

Gly Val Tyr Val Leu Pro Arg Arg Gly Pro Arg Leu Gly Val Arg Ala
35 40 45

Thr Arg Lys Thr Ser Glu Arg Ser Gln Pro Arg Gly Arg Arg Gln Pro
50 55 60

Ile Pro Lys Ala Arg Arg Ser Glu Gly Arg Ser Trp Ala Gln Pro Gly
65 70 75 80

Tyr Pro Trp Pro Leu Tyr Gly Asn Glu Gly Cys Gly Trp Ala Gly Trp
85 90 95

Leu Leu Ser Pro Arg Gly Ser Arg Pro Ser Trp Gly Pro Asn Asp Pro
100 105 110

Arg Arg Arg Ser Arg Asn Leu Gly Lys Val Ile Asp Thr Leu Thr Cys
115 120 125

Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Ala Pro Val
130 135 140

Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Ala Leu Glu Asp
145 150 155 160

Gly Ile Asn Phe Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe Ser Ile

165

170

175

Phe Leu Leu Ala Leu Phe Ser Cys Leu Val His Pro Ala Ala Ser
180 185 190

<210> 189

<211> 191

<212> PRT

<213> Homo sapiens

<220>

<223> Individual Isolate: S2

<400> 189

Met Ser Thr Leu Pro Lys Pro Gln Arg Lys Thr Lys Arg Asn Thr Ile
1 5 10 15

Arg Arg Pro Gln Asp Ile Lys Phe Pro Gly Gly Gln Ile Val Gly
20 25 30

Gly Val Tyr Val Leu Pro Arg Arg Gly Pro Arg Leu Gly Val Arg Ala
35 40 45

Thr Arg Lys Thr Ser Glu Arg Ser Gln Pro Arg Gly Arg Arg Gln Pro
50 55 60

Ile Pro Lys Ala Arg Arg Ser Glu Gly Arg Ser Trp Ala Gln Pro Gly
65 70 75 80

Tyr Pro Trp Pro Leu Tyr Gly Asn Glu Gly Cys Gly Trp Ala Gly Trp
85 90 95

Leu Leu Ser Pro Arg Gly Ser Arg Pro Ser Trp Gly Pro Asn Asp Pro
100 105 110

Arg Arg Arg Ser Arg Asn Leu Gly Lys Val Ile Asp Thr Leu Thr Cys
115 120 125

Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Ala Pro Val
130 135 140

Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Ala Leu Glu Asp
145 150 155 160

Gly Ile Asn Phe Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe Ser Ile
165 170 175

Phe Leu Leu Ala Leu Phe Ser Cys Leu Ile His Pro Ala Ala Ser
180 185 190

<210> 190
<211> 191
<212> PRT
<213> Homo sapiens

<220>
<223> Individual Isolate: DK12

<400> 190
Met Ser Thr Leu Pro Lys Pro Gln Arg Lys Thr Lys Arg Asn Thr Ile
1 5 10 15

Arg Arg Pro Gln Asp Val Lys Phe Pro Gly Gly Gln Ile Val Gly
20 25 30

Gly Val Tyr Val Leu Pro Arg Arg Gly Pro Arg Leu Gly Val Arg Ala
35 40 45

Thr Arg Lys Thr Ser Glu Arg Ser Gln Pro Arg Gly Arg Arg Gln Pro
50 55 60

Ile Pro Lys Ala Arg Arg Ser Glu Gly Arg Ser Trp Ala Gln Pro Gly
65 70 75 80

Tyr Pro Trp Pro Leu Tyr Gly Asn Glu Gly Cys Gly Trp Ala Gly Trp
85 90 95

Leu Leu Ser Pro Arg Gly Ser Arg Pro Ser Trp Gly Pro Asn Asp Pro
100 105 110

Arg Arg Arg Ser Arg Asn Leu Gly Lys Val Ile Asp Thr Leu Thr Cys
115 120 125

Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Ala Pro Val
130 135 140

Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Ala Leu Glu Asp
145 150 155 160

Gly Ile Asn Phe Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe Ser Ile
165 170 175

Phe Leu Leu Ala Leu Phe Ser Cys Leu Ile His Pro Ala Ala Ser
180 185 190

<210> 191
<211> 191
<212> PRT
<213> Homo sapiens

<220>
<223> Individual Isolate: Z4

<400> 191
Met Ser Thr Asn Pro Lys Pro Gln Arg Lys Thr Lys Arg Asn Thr Asn
1 5 10 15

Arg Arg Pro Met Asp Val Lys Phe Pro Gly Gly Gly Gln Ile Val Gly
20 25 30

Gly Val Tyr Leu Leu Pro Arg Arg Gly Pro Arg Leu Gly Val Arg Ala
35 40 45

Thr Arg Lys Thr Ser Glu Arg Ser Gln Pro Arg Gly Arg Arg Gln Pro
50 55 60

Ile Pro Lys Ala Arg Gln Pro Glu Gly Arg Ser Trp Ala Gln Pro Gly
65 70 75 80

Tyr Pro Trp Pro Leu Tyr Gly Asn Glu Gly Cys Gly Trp Ala Gly Trp
85 90 95

Leu Leu Ser Pro Arg Gly Ser Arg Pro Ser Trp Gly Pro Asn Asp Pro
100 105 110

Arg Arg Arg Ser Arg Asn Leu Gly Lys Val Ile Asp Thr Leu Thr Cys
115 120 125

Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Ile Val Gly Ala Pro Val
130 135 140

Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Ala Val Glu Asp
145 150 155 160

Gly Ile Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe Ser Ile
165 170 175

Phe Leu Leu Ala Leu Leu Ser Cys Leu Thr Val Pro Ala Ser Ala
180 185 190

<210> 192
<211> 191
<212> PRT
<213> Homo sapiens

<220>
<223> Individual Isolate: Z8

<400> 192
Met Ser Thr Asn Pro Lys Pro Gln Arg Lys Thr Lys Arg Asn Thr Asn
1 5 10 15

Arg Arg Pro Met Asp Val Lys Phe Pro Gly Gly Gln Ile Val Gly
20 25 30

Gly Val Tyr Leu Leu Pro Arg Arg Gly Pro Arg Leu Gly Val Arg Ala
35 40 45

Thr Arg Lys Thr Ser Glu Arg Ser Gln Pro Arg Gly Arg Arg Gln Pro
50 55 60

Ile Pro Lys Ala Arg Arg Ser Glu Gly Arg Ser Trp Ala Gln Pro Gly
65 70 75 80

Tyr Pro Trp Pro Leu Tyr Gly Asn Glu Gly Cys Gly Trp Ala Gly Trp
85 90 95

Leu Leu Ser Pro Arg Gly Ser Arg Pro Ser Trp Gly Pro Asn Asp Pro
100 105 110

Arg Arg Arg Ser Arg Asn Leu Gly Lys Val Ile Asp Thr Leu Thr Cys
115 120 125

Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Ala Pro Val
130 135 140

Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Ala Val Glu Asp
145 150 155 160

Gly Ile Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe Ser Ile
165 170 175

Phe Leu Leu Ala Leu Leu Ser Cys Leu Thr Val Pro Ala Ser Ala
180 185 190

<210> 193
<211> 191

<212> PRT

<213> Homo sapiens

<220>

<223> Individual Isolate: Z1

<400> 193

Met Ser Thr Asn Pro Lys Pro Gln Arg Lys Thr Lys Arg Asn Thr Asn
1 5 10 15

Arg Arg Pro Met Asp Val Lys Phe Pro Gly Gly Gln Ile Val Gly
20 25 30

Gly Val Tyr Leu Leu Pro Arg Arg Gly Pro Arg Leu Gly Val Arg Ala
35 40 45

Ala Arg Lys Thr Ser Glu Arg Ser Gln Pro Arg Gly Arg Arg Gln Pro
50 55 60

Ile Pro Lys Ala Arg Arg Ser Glu Gly Arg Ser Trp Ala Gln Pro Gly
65 70 75 80

Tyr Pro Trp Pro Leu Tyr Gly Asn Glu Gly Cys Gly Trp Ala Gly Trp
85 90 95

Leu Leu Ser Pro Arg Gly Ser Arg Pro Ser Trp Gly Pro Asn Asp Pro
100 105 110

Arg Arg Arg Ser Arg Asn Leu Gly Lys Val Ile Asp Thr Leu Thr Cys
115 120 125

Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Ala Pro Val
130 135 140

Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Ala Val Glu Asp
145 150 155 160

Gly Ile Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe Ser Ile
165 170 175

Phe Leu Leu Ala Leu Leu Ser Cys Leu Thr Thr Pro Ala Ser Ala
180 185 190

<210> 194

<211> 191

<212> PRT

<213> Homo sapiens

<220>

<223> Individual Isolate: Z5

<400> 194

Met Ser Thr Asn Pro Lys Pro Gln Arg Lys Thr Lys Arg Asn Thr Asn
1 5 10 15

Arg Arg Pro Met Asp Val Lys Phe Pro Gly Gly Gln Ile Val Gly
20 25 30

Gly Val Tyr Leu Leu Pro Arg Arg Gly Pro Arg Leu Gly Val Arg Ala
35 40 45

Thr Arg Lys Thr Ser Glu Arg Ser Gln Pro Arg Gly Arg Arg Gln Pro
50 55 60

Ile Pro Gln Ala Arg Arg Ser Glu Gly Arg Ser Trp Ala Gln Pro Gly
65 70 75 80

Tyr Pro Trp Pro Leu Tyr Gly Asn Glu Gly Cys Gly Trp Ala Gly Trp
85 90 95

Leu Leu Ser Pro Arg Gly Ser Arg Pro Ser Trp Gly Gln Asn Asp Pro
100 105 110

Arg Arg Arg Ser Arg Asn Leu Gly Lys Val Ile Asp Thr Leu Thr Cys
115 120 125

Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Ala Pro Val
130 135 140

Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Ala Leu Glu Asp
145 150 155 160

Gly Ile Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe Ser Ile
165 170 175

Phe Leu Leu Ala Leu Phe Ser Cys Leu Thr Thr Pro Ala Ser Ala
180 185 190

<210> 195

<211> 191

<212> PRT

<213> Homo sapiens

<220>

<223> Individual Isolate: Z6

<400> 195

Met Ser Thr Asn Pro Lys Pro Gln Arg Lys Thr Lys Arg Asn Thr Asn
1 5 10 15

Arg Arg Pro Met Asp Val Lys Phe Pro Gly Gly Gln Ile Val Gly
20 25 30

Gly Val Tyr Leu Leu Pro Arg Arg Gly Pro Arg Leu Gly Val Arg Ala
35 40 45

Thr Arg Lys Thr Ser Glu Arg Ser Gln Pro Arg Gly Arg Arg Gln Pro
50 55 60

Ile Pro Lys Ala Arg Arg Ser Glu Gly Arg Ser Trp Ala Gln Pro Gly
65 70 75 80

Tyr Pro Trp Pro Leu Tyr Gly Asn Glu Gly Cys Gly Trp Ala Gly Trp
85 90 95

Leu Leu Ser Pro Arg Gly Ser Arg Pro Ser Trp Gly Pro Asn Asp Pro
100 105 110

Arg Arg Arg Ser Arg Asn Leu Gly Lys Val Ile Asp Thr Leu Thr Cys
115 120 125

Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Ala Pro Val
130 135 140

Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Ala Val Glu Asp
145 150 155 160

Gly Ile Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe Ser Ile
165 170 175

Phe Leu Leu Ala Leu Leu Ser Cys Leu Thr Val Pro Thr Ser Ala
180 185 190

<210> 196

<211> 191

<212> PRT

<213> Homo sapiens

<220>

<223> Individual Isolate: Z7

<400> 196

Met Ser Thr Asn Pro Lys Pro Gln Arg Lys Thr Lys Arg Asn Thr Asn
1 5 10 15

Arg Arg Pro Met Asp Val Lys Phe Pro Gly Gly Gln Ile Val Gly
20 25 30

Gly Val Tyr Leu Leu Pro Arg Arg Gly Pro Arg Leu Gly Val Arg Thr
35 40 45

Thr Arg Lys Thr Ser Glu Arg Ser Gln Pro Arg Gly Arg Arg Gln Pro
50 55 60

Ile Pro Lys Ala Arg Arg Ser Glu Gly Arg Ser Trp Ala Gln Pro Gly
65 70 75 80

Tyr Pro Trp Pro Leu Tyr Gly Asn Glu Gly Cys Gly Trp Ala Gly Trp
85 90 95

Leu Leu Ser Pro Arg Gly Ser Arg Pro Ser Trp Gly Pro Asn Asp Pro
100 105 110

Arg Arg Arg Ser Arg Asn Leu Gly Lys Val Ile Asp Thr Leu Thr Cys
115 120 125

Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Ala Pro Val
130 135 140

Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Ala Leu Glu Asp
145 150 155 160

Gly Ile Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe Ser Ile
165 170 175

Phe Leu Leu Ala Leu Leu Ser Cys Leu Thr Val Pro Ala Ser Ala
180 185 190

<210> 197

<211> 191

<212> PRT

<213> Homo sapiens

<220>

<223> Individual Isolate: DK13

<400> 197

Met Ser Thr Asn Pro Lys Pro Gln Arg Lys Thr Lys Arg Asn Thr Asn

1	5	10	15
Arg Arg Pro Met Asp Val Lys Phe Pro Gly Gly Gly Gln Ile Val Gly			
20	25	30	
Gly Val Tyr Leu Leu Pro Arg Arg Gly Pro Arg Leu Gly Val Arg Ala			
35	40	45	
Thr Arg Lys Thr Ser Glu Arg Ser Gln Pro Arg Gly Arg Arg Gln Pro			
50	55	60	
Ile Pro Lys Ala Arg Gln Leu Glu Gly Arg Ser Trp Ala Gln Pro Gly			
65	70	75	80
Tyr Pro Trp Pro Leu Tyr Gly Asn Glu Gly Cys Gly Trp Ala Gly Trp			
85	90	95	
Leu Leu Ser Pro Arg Gly Ser Arg Pro Ser Trp Gly Pro Asn Asp Pro			
100	105	110	
Arg Arg Arg Ser Arg Asn Leu Gly Lys Val Ile Asp Thr Leu Thr Cys			
115	120	125	
Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Val Val Gly Ala Pro Val			
130	135	140	
Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Leu Leu Glu Asp			
145	150	155	160
Gly Val Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe Ser Ile			
165	170	175	
Phe Leu Leu Ala Leu Leu Ser Cys Leu Thr Val Pro Ala Ser Ala			
180	185	190	
<210> 198			
<211> 191			
<212> PRT			
<213> Homo sapiens			
<220>			
<223> Individual Isolate: SA4			
<400> 198			
Met Ser Thr Asn Pro Lys Pro Gln Arg Lys Thr Lys Arg Asn Thr Asn			
1	5	10	15

Arg Arg Pro Gln Asp Val Lys Phe Pro Gly Gly Gln Ile Val Gly
20 25 30

Gly Val Tyr Leu Leu Pro Arg Arg Gly Pro Arg Leu Gly Val Arg Ala
35 40 45

Thr Arg Lys Thr Ser Glu Arg Ser Gln Pro Arg Gly Arg Arg Gln Pro
50 55 60

Ile Pro Lys Ala Arg Gln Pro Thr Gly Arg Ser Trp Gly Gln Pro Gly
65 70 75 80

Tyr Pro Trp Pro Leu Tyr Ala Asn Glu Gly Leu Gly Trp Ala Gly Trp
85 90 95

Leu Leu Ser Pro Arg Gly Ser Arg Pro Asn Trp Gly Pro Asn Asp Pro
100 105 110

Arg Arg Lys Ser Arg Asn Leu Gly Lys Val Ile Asp Thr Leu Thr Cys
115 120 125

Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Pro Val
130 135 140

Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Val Leu Glu Asp
145 150 155 160

Gly Val Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe Ser Ile
165 170 175

Phe Ile Leu Ala Leu Leu Ser Cys Leu Thr Val Pro Ala Ser Ala
180 185 190

<210> 199

<211> 191

<212> PRT

<213> Homo sapiens

<220>

<223> Individual Isolate: SA5

<400> 199

Met Ser Thr Asn Pro Lys Pro Gln Arg Lys Thr Lys Arg Asn Thr Asn
1 5 10 15

Arg Arg Pro Gln Asp Val Lys Phe Pro Gly Gly Gln Ile Val Gly
20 25 30

Gly Val Tyr Leu Leu Pro Arg Arg Gly Pro Arg Leu Gly Val Arg Ala
35 40 45

Thr Arg Lys Thr Ser Glu Arg Ser Gln Pro Arg Gly Arg Arg Gln Pro
50 55 60

Ile Pro Lys Ala Arg Gln Pro Thr Gly Arg Ser Trp Gly Gln Pro Gly
65 70 75 80

Tyr Pro Trp Pro Leu Tyr Ala Asn Glu Gly Leu Gly Trp Ala Gly Trp
85 90 95

Leu Leu Ser Pro Arg Gly Ser Arg Pro Asn Trp Gly Pro Asn Asp Pro
100 105 110

Arg Arg Lys Ser Arg Asn Leu Gly Lys Val Ile Asp Thr Leu Thr Cys
115 120 125

Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Gly Pro Val
130 135 140

Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Val Leu Glu Asp
145 150 155 160

Gly Val Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe Ser Ile
165 170 175

Phe Ile Leu Ala Leu Leu Ser Cys Leu Thr Val Pro Ala Ser Ala
180 185 190

<210> 200

<211> 191

<212> PRT

<213> Homo sapiens

<220>

<223> Individual Isolate: SA7

<400> 200

Met Ser Thr Asn Pro Lys Pro Gln Arg Lys Thr Lys Arg Asn Thr Asn
1 5 10 15

Arg Arg Pro Gln Asp Val Lys Phe Pro Gly Gly Gln Ile Val Gly
20 25 30

Gly Val Tyr Leu Leu Pro Arg Arg Gly Pro Arg Leu Gly Val Arg Ala

35	40	45
Thr Arg Lys Thr Ser Glu Arg Ser Gln Pro Arg Gly Arg Arg Gln Pro		
50	55	60
Ile Pro Lys Ala Arg Gln Pro Thr Gly Arg Ser Trp Gly Gln Pro Gly		
65	70	75
Tyr Pro Trp Pro Leu Tyr Ala Asn Glu Gly Leu Gly Trp Ala Gly Trp		
85	90	95
Leu Leu Ser Pro Arg Gly Ser Arg Pro Asn Trp Gly Pro Asn Asp Pro		
100	105	110
Arg Arg Lys Ser Arg Asn Leu Gly Lys Val Ile Asp Thr Leu Thr Cys		
115	120	125
Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Pro Val		
130	135	140
Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Val Leu Glu Asp		
145	150	155
Gly Val Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe Ser Ile		
165	170	175
Phe Ile Leu Ala Leu Leu Ser Cys Leu Thr Val Pro Ala Ser Ala		
180	185	190
<210> 201		
<211> 191		
<212> PRT		
<213> Homo sapiens		
<220>		
<223> Individual Isolate: SA1		
<400> 201		
Met Ser Thr Asn Pro Lys Pro Gln Arg Lys Thr Lys Arg Asn Thr Asn		
1	5	10
15		
Leu Arg Pro Gln Asp Val Lys Phe Pro Gly Gly Gln Ile Val Gly		
20	25	30
Gly Val Tyr Leu Leu Pro Arg Arg Gly Pro Arg Leu Gly Val Arg Ala		
35	40	45

Thr Arg Lys Thr Ser Glu Arg Ser Gln Pro Arg Gly Arg Arg Gln Pro
50 55 60

Ile Pro Lys Ala Arg Gln Pro Thr Gly Arg Ser Trp Gly Gln Pro Gly
65 70 75 80

Tyr Pro Trp Pro Leu Tyr Ala Asn Glu Gly Leu Gly Trp Ala Gly Trp
85 90 95

Leu Leu Ser Pro Arg Gly Ser Arg Pro Asn Trp Gly Pro Asn Asp Pro
100 105 110

Arg Arg Lys Ser Arg Asn Leu Gly Lys Val Ile Asp Thr Leu Thr Cys
115 120 125

Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Gly Pro Val
130 135 140

Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Val Leu Glu Asp
145 150 155 160

Gly Val Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe Ser Ile
165 170 175

Phe Ile Leu Ala Leu Leu Ser Cys Leu Ile Ile Pro Ala Ser Ala
180 185 190

<210> 202

<211> 191

<212> PRT

<213> Homo sapiens

<220>

<223> Individual Isolate: SA3

<400> 202

Met Ser Thr Asn Pro Lys Pro Gln Arg Lys Thr Lys Arg Asn Thr Asn
1 5 10 15

Arg Arg Pro Gln Asp Val Lys Phe Pro Gly Gly Gln Ile Val Gly
20 25 30

Gly Val Tyr Leu Leu Pro Arg Arg Gly Pro Arg Leu Gly Val Arg Ala
35 40 45

Thr Arg Lys Thr Ser Glu Arg Ser Gln Pro Arg Gly Arg Arg Gln Pro
50 55 60

Ile Pro Lys Ala Arg Gln Pro Thr Gly Arg Ser Trp Gly Gln Pro Gly
65 70 75 80

Tyr Pro Trp Pro Leu Tyr Ala Asn Glu Gly Leu Glu Trp Ala Gly Trp
85 90 95

Leu Leu Ser Pro Arg Gly Ser Arg Pro Ser Trp Gly Pro Asn Asp Pro
100 105 110

Arg Arg Lys Ser Arg Asn Leu Gly Lys Val Ile Asp Thr Leu Thr Cys
115 120 125

Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Gly Pro Val
130 135 140

Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Val Leu Glu Asp
145 150 155 160

Gly Val Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe Ser Ile
165 170 175

Phe Ile Leu Ala Leu Leu Ser Cys Leu Thr Val Pro Ala Ser Ala
180 185 190

<210> 203

<211> 191

<212> PRT

<213> Homo sapiens

<220>

<223> Individual Isolate: SA13

<400> 203

Met Ser Thr Asn Pro Lys Pro Gln Arg Lys Thr Lys Arg Asn Thr Asn
1 5 10 15

Arg Arg Pro Gln Asp Val Lys Phe Pro Gly Gly Gln Ile Val Gly
20 25 30

Gly Val Tyr Leu Leu Pro Arg Arg Gly Pro Arg Leu Gly Val Arg Ala
35 40 45

Thr Arg Lys Thr Ser Glu Arg Ser Gln Pro Arg Gly Arg Arg Gln Pro
50 55 60

Ile Pro Lys Ala Arg Gln Pro Thr Gly Arg Ser Trp Gly Gln Pro Gly

65 70 75 80

Tyr Pro Trp Pro Leu Tyr Ala Asn Glu Gly Leu Gly Trp Ala Gly Trp
85 90 95

Leu Leu Ser Pro Arg Gly Ser Arg Pro Asn Trp Gly Pro Asn Asp Pro
100 105 110

Arg Arg Lys Ser Arg Asn Leu Gly Lys Val Ile Asp Thr Leu Thr Cys
115 120 125

Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Gly Pro Val
130 135 140

Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Val Leu Glu Asp
145 150 155 160

Gly Val Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe Ser Ile
165 170 175

Phe Ile Leu Ala Leu Leu Ser Cys Leu Thr Val Pro Thr Ser Ala
180 185 190

<210> 204

<211> 191

<212> PRT

<213> Homo sapiens

<220>

<223> Individual Isolate: SA6

<400> 204

Met Ser Thr Asn Pro Lys Pro Gln Arg Lys Thr Gln Arg Asn Thr Asn
1 5 10 15

Arg Arg Pro Gln Asp Val Lys Phe Pro Gly Gly Gln Ile Val Gly
20 25 30

Gly Val Tyr Leu Leu Pro Arg Arg Gly Pro Arg Met Gly Val Arg Ala
35 40 45

Thr Arg Lys Thr Ser Glu Arg Ser Gln Pro Arg Gly Arg Arg Gln Pro
50 55 60

Ile Pro Lys Ala Arg Gln Ser Ala Gly Arg Ser Trp Gly Gln Pro Gly
65 70 75 80

Tyr Pro Trp Pro Leu Tyr Ala Asn Glu Gly Leu Gly Trp Ala Gly Trp
85 90 95

Leu Leu Ser Pro Arg Gly Ser Arg Pro Asn Trp Gly Pro Asn Asp Pro
100 105 110

Arg Arg Lys Ser Arg Asn Leu Gly Lys Val Ile Asp Thr Leu Thr Cys
115 120 125

Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Gly Pro Val
130 135 140

Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Val Leu Glu Asp
145 150 155 160

Gly Val Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe Ser Ile
165 170 175

Phe Val Leu Ala Leu Leu Ser Cys Leu Thr Val Pro Ala Ser Ala
180 185 190

<210> 205

<211> 191

<212> PRT

<213> Homo sapiens

<220>

<223> Individual Isolate: SA11

<400> 205

Met Ser Thr Asn Pro Lys Pro Gln Arg Lys Thr Lys Arg Asn Thr Asn
1 5 10 15

Arg Arg Pro Gln Asp Val Lys Phe Pro Gly Gly Gln Ile Val Gly
20 25 30

Gly Val Tyr Leu Leu Pro Arg Arg Gly Pro Arg Leu Gly Val Arg Ala
35 40 45

Thr Arg Lys Thr Ser Glu Arg Ser Gln Pro Arg Gly Arg Arg Gln Pro
50 55 60

Ile Pro Lys Ala Arg Gln Pro Thr Gly Arg Ser Trp Gly Gln Pro Gly
65 70 75 80

Tyr Pro Trp Pro Phe Tyr Ala Asn Glu Gly Leu Gly Trp Ala Gly Trp
85 90 95

Leu Leu Ser Pro Arg Gly Ser Arg Pro Asn Trp Gly Pro Asn Asp Pro
100 105 110

Arg Arg Arg Ser Arg Asn Leu Gly Lys Val Ile Asp Thr Leu Thr Cys
115 120 125

Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Gly Pro Val
130 135 140

Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Ala Leu Glu Asp
145 150 155 160

Gly Val Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe Ser Ile
165 170 175

Phe Ile Leu Ala Leu Leu Ser Cys Leu Thr Val Pro Ala Thr Ala
180 185 190

<210> 206

<211> 191

<212> PRT

<213> Homo sapiens

<220>

<223> Individual Isolate: HK2

<400> 206

Met Ser Thr Leu Pro Lys Pro Gln Arg Lys Thr Lys Arg Asn Thr Asn
1 5 10 15

Arg Arg Pro Thr Asp Val Lys Phe Pro Gly Gly Gln Ile Val Gly
20 25 30

Gly Val Tyr Leu Leu Pro Arg Arg Gly Pro Arg Leu Gly Val Arg Ala
35 40 45

Thr Arg Lys Thr Ser Glu Arg Ser Gln Pro Arg Gly Arg Arg Gln Pro
50 55 60

Ile Pro Lys Ala Arg Gln Pro Gln Gly Arg His Trp Ala Gln Pro Gly
65 70 75 80

Tyr Pro Trp Pro Leu Tyr Gly Asn Glu Gly Cys Gly Trp Ala Gly Trp
85 90 95

Leu Leu Ser Pro Arg Gly Ser Arg Pro His Trp Gly Pro Asn Asp Pro

100	105	110
Arg Arg Arg Ser Arg Asn Leu Gly Lys Val Ile Asp Thr Leu Thr Cys		
115	120	125
Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Val Val Gly Ala Pro Leu		
130	135	140
Gly Gly Val Ala Ala Ala Leu Ala His Gly Val Arg Ala Ile Glu Asp		
145	150	155
Gly Ile Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe Ser Ile		
165	170	175
Phe Leu Leu Ala Leu Leu Ser Cys Leu Thr Thr Pro Ala Ser Ala		
180	185	190

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42

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aatggcaccy tgcrcgtctg gataacaagtr acacctaattg tggctgtgaa acac	54
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tgarctagyc ctysargtyg tcttcgggygg y	31

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<211> 42

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<213> Homo sapiens

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<210> 231

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atctagcatc ttgagggtagt ac ctgagatttg tgcgagtgtg atatttggtg gc

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<222> (26)
<223> "Val" or "Met"

<400> 240

Trp Ile Gln Val Thr Pro Asn Val Ala Val Lys His Arg Gly Ala Leu
1 5 10 15

Thr His Asn Leu Arg Xaa His Xaa Asp Xaa Ile Val Met Ala Ala Thr
20 25 30

Val

<210> 241

<211> 33

<212> PRT

<213> Homo sapiens

<400> 241

Trp Val Pro Val Ala Pro Asn Leu Ala Ile Ser Gln Pro Gly Ala Leu
1 5 10 15

Thr Lys Gly Leu Arg Ala His Ile Asp Ile Ile Val Met Ser Ala Thr
20 25 30

Val

<210> 242

<211> 33

<212> PRT

<213> Homo sapiens

<220>

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<222> (11)

<223> "Arg" or "Gln"

<220>

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<222> (12)

<223> "Arg" or "Gln"

<400> 242

Trp Ile Pro Val Xaa Pro Asn Val Ala Val Xaa Xaa Pro Gly Ala Leu
1 5 10 15

Thr Gln Gly Leu Arg Thr His Ile Asp Met Val Val Met Ser Ala Thr
20 25 30

Leu

<210> 243
<211> 33
<212> PRT
<213> Homo sapiens

<220>
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<220>
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<400> 243
Trp Thr Xaa Val Thr Pro Thr Val Ala Val Arg Tyr Val Gly Ala Thr
1 5 10 15

Thr Ala Ser Ile Arg Ser His Val Asp Leu Leu Val Gly Ala Ala Thr
20 25 30

Xaa

<210> 244
<211> 33
<212> PRT
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<220>
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Trp Val Ala Leu Xaa Pro Thr Leu Ala Ala Arg Asn Xaa Xaa Xaa
1 5 10 15

Thr Xaa Xaa Ile Arg Xaa His Val Asp Leu Leu Val Gly Ala Ala Xaa
20 25 30

Phe

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<210> 245
<211> 33
<212> PRT
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<400> 245
Trp Val Xaa Xaa Xaa Pro Thr Val Ala Thr Arg Asp Gly Lys Leu Pro
1 5 10 15

Xaa Xaa Gln Leu Arg Arg Xaa Ile Asp Leu Leu Val Gly Ser Ala Thr
20 25 30

Leu

<210> 246

<211> 33
<212> PRT
<213> Homo sapiens

<400> 246
Trp Thr Pro Val Thr Pro Thr Val Ala Val Ala His Pro Gly Ala Pro
1 5 10 15

Leu Glu Ser Phe Arg Arg His Val Asp Leu Met Val Gly Ala Ala Thr
20 25 30

Leu

<210> 247
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<400> 247
Trp Val Ala Leu Thr Pro Thr Val Ala Xaa Xaa Tyr Ile Gly Ala Pro
1 5 10 15

Leu Xaa Ser Xaa Arg Arg His Val Asp Leu Met Val Gly Ala Ala Thr
20 25 30

Val

<210> 248
<211> 33
<212> PRT
<213> Homo sapiens

<400> 248
Trp Val Ser Leu Thr Pro Thr Val Ala Ala Gln His Leu Asn Ala Pro
1 5 10 15

Leu Glu Ser Leu Arg Arg His Val Asp Leu Met Val Gly Gly Ala Thr
20 25 30

Leu

<210> 249
<211> 33
<212> PRT
<213> Homo sapiens

<400> 249
Trp Val Pro Leu Thr Pro Thr Val Ala Ala Pro Tyr Pro Asn Ala Pro
1 5 10 15

Leu Glu Ser Met Arg Arg His Val Asp Leu Met Val Gly Ala Ala Thr
20 25 30

Met

<210> 250
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<212> PRT
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Trp Val Xaa Ile Thr Pro Thr Leu Ser Ala Pro Xaa Xaa Gly Ala Val
1 5 10 15

Thr Ala Pro Leu Arg Arg Xaa Val Asp Tyr Leu Ala Gly Gly Ala Ala
20 25 30

Leu

<210> 251

<211> 33

<212> PRT

<213> Homo sapiens

<400> 251

Trp His Ala Val Thr Pro Thr Leu Ala Ile Pro Asn Ala Ser Thr Pro
1 5 10 15

Ala Thr Gly Phe Arg Arg His Val Asp Leu Leu Ala Gly Ala Ala Val
20 25 30

Val

<210> 252

<211> 23

<212> PRT

<213> Homo sapiens

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<223> "Glu" or "Gln"

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Thr Leu Thr Met Ile Leu Ala Tyr Ala Ala Arg Val Pro Glu Leu Xaa
1 5 10 15

Leu Xaa Val Val Phe Gly Gly
20

<210> 253
<211> 23
<212> PRT
<213> Homo sapiens

<400> 253
Thr Thr Thr Met Leu Leu Ala Tyr Leu Val Arg Ile Pro Glu Val Ile
1 5 10 15

Leu Asp Ile Val Thr Gly Gly
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Thr Xaa Thr Xaa Ile Leu Ala Tyr Xaa Met Arg Val Pro Glu Val Ile

1

5

10

15

Xaa Asp Ile Xaa Xaa Gly Ala

20

<210> 255

<211> 23

<212> PRT

<213> Homo sapiens

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<223> "Leu" or "Val"

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<223> "Ile" or "Leu"

<400> 255

Ala Val Gly Met Val Val Ala His Xaa Leu Arg Leu Pro Gln Thr Xaa

1

5

10

15

Phe Asp Ile Xaa Ala Gly Ala

20

<210> 256
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<400> 256
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1 5 10 15

Xaa Asp Xaa Val Xaa Gly Ala
20

<210> 257
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Thr Xaa Ala Leu Val Xaa Ala Gln Leu Leu Arg Xaa Pro Gln Ala Xaa
1 5 10 15

Leu Asp Met Ile Ala Gly Ala
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<210> 258
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<212> PRT
<213> Homo sapiens

<400> 258
Thr Thr Thr Leu Leu Ala Gln Ile Met Arg Val Pro Thr Ala Phe
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Leu Asp Met Val Ala Gly Gly
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1 5 10 15

Val Asp Leu Leu Xaa Gly Gly
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<210> 260
<211> 23
<212> PRT
<213> Homo sapiens

<400> 260
Thr Ala Thr Leu Val Leu Ala Gln Leu Met Arg Ile Pro Gly Ala Met
1 5 10 15

Val Asp Leu Leu Ala Gly Gly
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<210> 261
<211> 23
<212> PRT
<213> Homo sapiens

<400> 261
Thr Ser Ala Leu Ile Met Ala Gln Ile Leu Arg Ile Pro Ser Ile Leu
1 5 10 15

Gly Asp Leu Leu Thr Gly Gly
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<210> 262
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<223> "Gly" or "Ala"

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1 5 10 15
Ile Asp Ile Ile Ala Gly Xaa
20

<210> 263
<211> 23
<212> PRT
<213> Homo sapiens

<400> 263
Thr Thr Thr Leu Val Leu Ser Ser Ile Leu Arg Val Pro Glu Ile Cys
1 5 10 15
Ala Ser Val Ile Phe Gly Gly
20

<210> 264

<211> 191
<212> PRT
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<223> "Thr" or "Pro"

<400> 264

Met Ser Thr Asn Pro Lys Pro Gln Arg Lys Thr Lys Arg Asn Thr Asn
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Arg Arg Pro Gln Asp Val Lys Phe Pro Gly Gly Gln Ile Val Gly
20 25 30

Gly Val Tyr Leu Leu Pro Arg Arg Gly Pro Arg Leu Gly Val Arg Ala
35 40 45

Xaa Arg Lys Thr Ser Glu Arg Ser Gln Pro Arg Gly Arg Arg Gln Pro
50 55 60

Ile Pro Lys Ala Arg Arg Pro Glu Gly Arg Thr Trp Ala Gln Pro Gly
65 70 75 80

Tyr Pro Trp Pro Leu Tyr Gly Asn Glu Gly Cys Gly Trp Ala Gly Trp
85 90 95

Leu Leu Ser Pro Arg Gly Ser Arg Pro Ser Trp Gly Pro Thr Asp Pro
100 105 110

Arg Arg Arg Ser Arg Asn Leu Gly Lys Val Ile Asp Thr Leu Thr Cys
115 120 125

Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Ala Pro Leu
130 135 140

Gly Gly Ala Ala Arg Ala Leu Ala His Gly Val Arg Val Leu Glu Asp
145 150 155 160

Gly Val Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe Ser Ile
165 170 175

Phe Leu Leu Ala Leu Leu Ser Cys Leu Thr Val Pro Ala Ser Ala
180 185 190

<210> 265

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<223> "Ala" or "Val"

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Met Ser Thr Xaa Pro Lys Pro Gln Arg Xaa Thr Lys Arg Asn Thr Xaa
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Arg Arg Pro Gln Asp Val Lys Phe Pro Gly Gly Gln Ile Val Gly
20 25 30

Gly Val Tyr Leu Leu Pro Arg Arg Gly Pro Arg Leu Gly Val Arg Ala
35 40 45

Thr Arg Lys Thr Ser Glu Arg Ser Gln Pro Arg Gly Arg Arg Gln Pro
50 55 60

Ile Pro Lys Ala Arg Xaa Pro Glu Gly Arg Xaa Trp Ala Gln Pro Gly
65 70 75 80

Xaa Pro Trp Pro Leu Tyr Xaa Xaa Glu Gly Xaa Gly Trp Ala Gly Trp
85 90 95

Leu Leu Ser Pro Xaa Gly Ser Arg Pro Xaa Trp Gly Pro Xaa Asp Pro
100 105 110

Arg Arg Arg Ser Arg Asn Leu Gly Lys Val Ile Asp Thr Leu Thr Cys
115 120 125

Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Xaa Pro Leu
130 135 140

Gly Gly Xaa Ala Arg Ala Leu Ala His Gly Val Arg Val Xaa Glu Asp
145 150 155 160

Gly Val Asn Tyr Ala Thr Gly Asn Xaa Pro Gly Cys Xaa Phe Ser Ile
165 170 175

Phe Leu Leu Ala Leu Leu Ser Cys Leu Thr Xaa Pro Xaa Ser Ala
180 185 190

<210> 266

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<213> Homo sapiens

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<223> "Val" or "Ala"

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Met Ser Thr Xaa Pro Lys Pro Gln Arg Xaa Thr Lys Arg Asn Thr Xaa
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Arg Arg Pro Gln Asp Val Lys Phe Pro Gly Gly Gln Ile Val Gly
20 25 30

Gly Val Tyr Leu Leu Pro Arg Arg Gly Pro Arg Leu Gly Val Arg Ala
35 40 45

Xaa Arg Lys Thr Ser Glu Arg Ser Gln Pro Arg Gly Arg Arg Gln Pro
50 55 60

Ile Pro Lys Ala Arg Xaa Pro Glu Gly Arg Xaa Trp Ala Gln Pro Gly
65 70 75 80

Xaa Pro Trp Pro Leu Tyr Xaa Xaa Glu Gly Xaa Gly Trp Ala Gly Trp
85 90 95

Leu Leu Ser Pro Xaa Gly Ser Arg Pro Xaa Trp Gly Pro Xaa Asp Pro
100 105 110

Arg Arg Arg Ser Arg Asn Leu Gly Lys Val Ile Asp Thr Leu Thr Cys
115 120 125

Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Xaa Pro Leu
130 135 140

Gly Gly Xaa Ala Arg Ala Leu Ala His Gly Val Arg Val Xaa Glu Asp
145 150 155 160

Gly Val Asn Tyr Ala Thr Gly Asn Xaa Pro Gly Cys Xaa Phe Ser Ile
165 170 175

Phe Leu Leu Ala Leu Leu Ser Cys Leu Thr Xaa Pro Xaa Ser Ala
180 185 190

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<223> "Thr" or "Ala"

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<223> "Phe" or "Leu"

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<223> "Ala" or "Val"

<400> 267

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1 5 10 15

Arg Arg Pro Gln Asp Val Lys Phe Pro Gly Gly Gln Ile Val Gly
20 25 30

Gly Val Tyr Leu Leu Pro Arg Arg Gly Pro Arg Leu Gly Val Arg Xaa
35 40 45

Thr Arg Lys Thr Ser Glu Arg Ser Gln Pro Arg Gly Arg Arg Gln Pro
50 55 60

Ile Pro Lys Asp Arg Arg Xaa Thr Gly Lys Ser Trp Gly Lys Pro Gly
65 70 75 80

Tyr Pro Trp Pro Leu Tyr Gly Asn Glu Gly Leu Gly Trp Ala Gly Trp
85 90 95

Leu Leu Ser Pro Arg Gly Ser Arg Pro Ser Trp Gly Pro Xaa Asp Pro
100 105 110

Arg His Arg Ser Arg Asn Val Gly Lys Val Ile Asp Thr Leu Thr Cys
115 120 125

Xaa Xaa Ala Asp Leu Met Gly Tyr Xaa Pro Val Val Gly Xaa Pro Leu
130 135 140

Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Val Leu Glu Asp
145 150 155 160

Gly Val Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe Ser Ile
165 170 175

Phe Leu Leu Ala Leu Leu Ser Cys Ile Thr Xaa Pro Xaa Ser Ala
180 185 190

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<223> "Cys" or "Phe" or "Ala"

<400> 268

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1

5

10

15

Arg Arg Pro Gln Asp Val Lys Phe Pro Gly Gly Gln Ile Val Gly

20

25

30

Gly Val Tyr Leu Leu Pro Arg Arg Gly Pro Arg Leu Gly Val Arg Xaa

35

40

45

Thr	Arg	Lys	Xaa	Ser	Glu	Arg	Ser	Gln	Pro	Arg	Gly	Arg	Arg	Gln	Pro
50				55				60							
Ile	Pro	Lys	Asp	Arg	Arg	Ser	Thr	Gly	Lys	Xaa	Trp	Gly	Lys	Pro	Gly
65				70				75				80			
Tyr	Pro	Trp	Pro	Leu	Tyr	Gly	Asn	Glu	Gly	Cys	Gly	Trp	Ala	Gly	Trp
				85				90				95			
Leu	Leu	Ser	Pro	Arg	Gly	Ser	Xaa	Pro	Xaa	Trp	Gly	Pro	Thr	Asp	Pro
				100				105				110			
Arg	His	Xaa	Ser	Arg	Asn	Leu	Gly	Xaa	Val	Ile	Asp	Thr	Ile	Thr	Cys
				115				120			125				
Gly	Phe	Ala	Asp	Leu	Met	Gly	Tyr	Ile	Pro	Val	Val	Gly	Ala	Pro	Val
				130				135			140				
Gly	Gly	Val	Ala	Arg	Ala	Leu	Ala	His	Gly	Val	Arg	Val	Leu	Glu	Asp
				145				150			155			160	
Gly	Ile	Asn	Tyr	Ala	Thr	Gly	Asn	Leu	Pro	Gly	Cys	Ser	Phe	Ser	Ile
				165				170				175			
Phe	Leu	Leu	Ala	Leu	Leu	Ser	Cys	Xaa	Thr	Val	Pro	Val	Ser	Ala	
				180				185			190				

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<223> "Val" or "Ala"

<400> 269
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Arg Arg Pro Gln Asp Val Lys Phe Pro Gly Gly Gln Ile Val Gly
20 25 30

Gly Val Tyr Leu Leu Pro Arg Arg Gly Pro Arg Leu Gly Val Arg Xaa
35 40 45

Thr Arg Lys Xaa Ser Glu Arg Ser Gln Pro Arg Gly Arg Arg Gln Pro
50 55 60

Ile Pro Lys Asp Arg Arg Xaa Thr Gly Lys Xaa Trp Gly Xaa Pro Gly
65 70 75 80

Tyr Pro Trp Pro Leu Tyr Gly Asn Glu Gly Xaa Gly Trp Ala Gly Trp
85 90 95

Leu Leu Ser Pro Arg Gly Ser Xaa Pro Xaa Trp Gly Pro Xaa Asp Pro
100 105 110

Arg His Xaa Ser Arg Asn Xaa Gly Xaa Val Ile Asp Thr Xaa Thr Cys
115 120 125

Xaa Xaa Ala Asp Leu Met Gly Tyr Xaa Pro Val Val Gly Xaa Pro Xaa
130 135 140

Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Val Leu Glu Asp
145 150 155 160

Gly Xaa Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe Ser Ile
165 170 175

Phe Leu Leu Ala Leu Leu Ser Cys Xaa Xaa Xaa Pro Xaa Ser Ala
180 185 190

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<223> "Ile" or "Val"

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Arg Arg Pro Gln Asp Xaa Lys Phe Pro Gly Gly Gly Gln Ile Val Gly
20 25 30

Gly Val Tyr Val Leu Pro Arg Arg Gly Pro Arg Leu Gly Val Arg Ala
35 40 45

Thr Arg Lys Thr Ser Glu Arg Ser Gln Pro Arg Gly Arg Arg Gln Pro
50 55 60

Ile Pro Lys Ala Arg Arg Ser Glu Gly Arg Ser Trp Ala Gln Pro Gly
65 70 75 80

Tyr Pro Trp Pro Leu Tyr Gly Asn Glu Gly Cys Gly Trp Ala Gly Trp
85 90 95

Leu Leu Ser Pro Arg Gly Ser Arg Pro Ser Trp Gly Pro Asn Asp Pro
100 105 110

Arg Arg Arg Ser Arg Asn Leu Gly Lys Val Ile Asp Thr Leu Thr Cys
115 120 125

Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Ala Pro Val
130 135 140

Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Ala Leu Glu Asp
145 150 155 160

Gly Ile Asn Phe Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe Ser Ile
165 170 175

Phe Leu Leu Ala Leu Phe Ser Cys Leu Xaa His Pro Ala Ala Ser
180 185 190

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<223> "Ala" or "Thr"

<400> 271

Met Ser Thr Asn Pro Lys Pro Gln Arg Lys Thr Lys Arg Asn Thr Asn
1 5 10 15

Arg Arg Pro Met Asp Val Lys Phe Pro Gly Gly Gln Ile Val Gly
20 25 30

Gly Val Tyr Leu Leu Pro Arg Arg Gly Pro Arg Leu Gly Val Arg Xaa
35 40 45

Xaa Arg Lys Thr Ser Glu Arg Ser Gln Pro Arg Gly Arg Arg Gln Pro
50 55 60

Ile Pro Xaa Ala Arg Xaa Xaa Glu Gly Arg Ser Trp Ala Gln Pro Gly
65 70 75 80

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Xaa Arg Pro Gln Asp Val Lys Phe Pro Gly Gly Gln Ile Val Gly
20 25 30

Gly Val Tyr Leu Leu Pro Arg Arg Gly Pro Arg Xaa Gly Val Arg Ala
35 40 45

Thr Arg Lys Thr Ser Glu Arg Ser Gln Pro Arg Gly Arg Arg Gln Pro
50 55 60

Ile Pro Lys Ala Arg Gln Xaa Xaa Gly Arg Ser Trp Gly Gln Pro Gly
65 70 75 80

Tyr Pro Trp Pro Xaa Tyr Ala Asn Glu Gly Leu Xaa Trp Ala Gly Trp
85 90 95

Leu Leu Ser Pro Arg Gly Ser Arg Pro Xaa Trp Gly Pro Asn Asp Pro
100 105 110

Arg Arg Xaa Ser Arg Asn Leu Gly Lys Val Ile Asp Thr Leu Thr Cys
115 120 125

Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Pro Val
130 135 140

Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Xaa Leu Glu Asp
145 150 155 160

Gly Val Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe Ser Ile
165 170 175

Phe Xaa Leu Ala Leu Leu Ser Cys Leu Xaa Xaa Pro Xaa Xaa Ala
180 185 190

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Gly Val Tyr Xaa Leu Pro Arg Arg Gly Pro Arg Xaa Gly Val Arg Xaa
35 40 45

Xaa Arg Lys Xaa Ser Glu Arg Ser Gln Pro Arg Gly Arg Arg Gln Pro
50 55 60

Ile Pro Xaa Xaa Arg Xaa Xaa Xaa Gly Xaa Xaa Trp Xaa Xaa Pro Gly
65 70 75 80

Xaa Pro Trp Pro Xaa Tyr Xaa Xaa Glu Gly Xaa Xaa Trp Ala Gly Trp
85 90 95

Leu Leu Ser Pro Xaa Gly Ser Xaa Pro Xaa Trp Gly Xaa Xaa Asp Pro
100 105 110

Arg Xaa Xaa Ser Arg Asn Xaa Gly Xaa Val Ile Asp Thr Xaa Thr Cys
115 120 125

Xaa Xaa Ala Asp Leu Met Gly Tyr Xaa Pro Xaa Val Gly Xaa Pro Xaa
130 135 140

Gly Gly Xaa Ala Xaa Ala Leu Ala His Gly Val Arg Xaa Xaa Glu Asp
145 150 155 160

Gly Xaa Asn Xaa Ala Thr Gly Asn Xaa Pro Gly Cys Xaa Phe Ser Ile
165 170 175

Phe Xaa Leu Ala Leu Xaa Ser Cys Xaa Xaa Xaa Pro Xaa Xaa Xaa
180 185 190

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<212> PRT
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Xaa Arg Pro Xaa Asp Xaa Lys Phe Pro Gly Gly Gly Gln Ile Val Gly															
20 25 30															
Gly Val Tyr Xaa Leu Pro Arg Arg Gly Pro Arg Xaa Gly Val Arg Xaa															
35 40 45															
Xaa Arg Lys Xaa Ser Glu Arg Ser Gln Pro Arg Gly Arg Arg Gln Pro															
50 55 60															
Ile Pro Xaa Xaa Arg Xaa Xaa Xaa Gly Xaa Xaa Trp Xaa Xaa Pro Gly															
65 70 75 80															
Xaa Pro Trp Pro Xaa Tyr Xaa Xaa Glu Gly Xaa Xaa Trp Ala Gly Trp															
85 90 95															
Leu Leu Ser Pro Xaa Gly Ser Xaa Pro Xaa Trp Gly Xaa Xaa Asp Pro															
100 105 110															
Arg Xaa Xaa Ser Arg Asn Xaa Gly Xaa Val Ile Asp Thr Xaa Thr Cys															
115 120 125															
Xaa Xaa Ala Asp Leu Met Gly Tyr Xaa Pro Xaa Val Gly Xaa Pro Xaa															
130 135 140															
Gly Gly Xaa Ala Xaa Ala Leu Ala His Gly Val Arg Xaa Xaa Glu Asp															
145 150 155 160															
Gly Xaa Asn Xaa Ala Thr Gly Asn Xaa Pro Gly Cys Xaa Phe Ser Ile															
165 170 175															
Phe Xaa Leu Ala Leu Xaa Ser Cys Xaa Xaa Xaa Pro Xaa Xaa Xaa															
180 185 190															